

Box seg listing /1623

ATTORNEY DOCKET NUMBER: 00786/351005

Certificate of Mailing:	Date of Denosit:	August 1, 2001
Cerunicale of Maning.	Date of Deposit.	

I hereby certify under 37 C.F.R. § 1.8(a) that this correspondence is being deposited with the United States Postal Service as first class mail with sufficient postage on the date indicated above and is addressed to the Assistant Commissioner of Patents, Washington, D.C. 20231.

Colleen Coyne

Printed name of person mailing correspondence

Signature of person mailin

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Applicant:

Gary Ruvkun et al.

Art Unit:

1633

Serial No.:

09/844,353

Examiner:

Filed:

April 27, 2001

Customer No.:

21559

Title:

THERAPEUTIC AND DIAGNOSTIC TOOLS FOR IMPAIRED

GLUCOSE TOLERANCE CONDITIONS

Assistant Commissioner For Patents

Washington, D.C. 20231

STATEMENT UNDER 37 C.F.R. § 1.825

In reply to the Notice to Comply mailed June 4, 2001 and as required by 37 C.F.R. § 1.825(a), enclosed is an amended sequence listing consisting of 63 sheets to be inserted at the end of the application.

The amendments correct typographical errors. I hereby submit that the substitute sheets contain no new matter.

As required by 37 C.F.R. § 1.825(b), enclosed is a diskette containing a copy of the sequence listing in computer readable form including all previously submitted data with the amendments incorporated therein. The contents of the computer readable form are the same as the contents of the paper sheets.

If there are any charges or any credits, please apply them to Deposit Account No.

03-2095.

Respectfully submitted,

Date: 1 August 2001

Karen L. Elbing, Ph.D.

Reg. No. 35,238

Clark & Elbing LLP 176 Federal Street Boston, MA 02110

Telephone: 617-428-0200 Facsimile: 617-428-7045

21559



SEQUENCE LISTING

RECEIVED

AUG 0 8 2001

TECH CENTER 1600/2900

20

18

<110> Ruvkun, Gary Kimura, Koutarou Patterson, Garth Ogg, Scott Paradis, Suzanne Tissenbaum, Heidi Morris, Jason Koweek, Allison

<120> THERAPEUTIC AND DIAGNOSTIC TOOLS FOR IMPAIRED GLUCOSE TOLERANCE CONDITIONS

<130> 00786/351005

<140> US 09/844,353

<141> 2001-04-27

<150> US 08/857,076

<151> 1997-05-15

<160> 114

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer/probe derived from C. elegans

<400> 1

cgctacggca aaaaagtgaa

<210> 2

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer/probe derived from C. elegans

<400> 2

cgatgatgaa gatacccc

<210> 3

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer/probe derived from C. elegans

<400> 3 tgatgegaac ggcgatcgat	20
<210> 4 <211> 20 <212> DNA <213> Artificial Sequence	
<220> <223> Primer/probe derived from C. elegans	
<400> 4 acgctggatc atctacatta	20
<210> 5 <211> 22 <212> DNA <213> Artificial Sequence	
<220> <223> Primer/probe derived from C. elegans	
<400> 5 ggtttaatta cccaagtttg ag	22
<210> 6 <211> 20 <212> DNA <213> Artificial Sequence	
<220> <223> Primer/probe derived from C. elegans	
<400> 6 gctcacgggt cacacaacga	20
<210> 7 <211> 20 <212> DNA <213> Artificial Sequence	
<220> <223> Primer/probe derived from C. elegans	
<400> 7 tgatgcgaac ggcgatcgat	20
<210> 8 <211> 21 <212> DNA <213> Artificial Sequence	
<220> <223> Primer/probe derived from C. elegans	
<400> 8 tgagggccaa ctaaagaaga c	21

```
<210> 9
<211> 20
<212> DNA
<213> Artificial Sequence
<223> Primer/probe derived from C. elegans
<400> 9
                                                                   20
cgctacggca aaaaagtgaa
<210> 10
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> Primer/probe derived from C. elegans
<400> 10
                                                                   20
gacgatcccg aggtgagtat
<210> 11
<211> 5816
<212> DNA
<213> Caenorhabditis elegans
<220>
<221> misc_feature
<222> (1)...(5816)
<223> n = A, T, C or G
<400> 11
ggtttaatta cccaagtttg agctccaaga gcacacatct gatcgtcgga ttctactgta 60
ctccccgaaa aaccaacaaa aaacacaagt ttttgaacac ttgtaaatgc agacagaacg 120
atgacgagaa tgaatattgt cagatgtcgg agacgacaca aaattttgga aaatttggaa 180
gaagagaatc teggeeegag etgetegteg aegaetteaa caacegetge caeegaaget 240
ctcggaacaa ccactgagga tatgaggctt aagcagcagc gaagctcgtc gcgtgccacg 300
gagcacgata ttgtcgacgg caatcaccac gacgacgagc acatcacaat gagacggctt 360
cgacttgtca aaaattcgcg gacgcggcgt agaacgacgc ccgattcaag tatggactgc 420
tatgaggaaa acccgccatc acaaaaactt caataaatta ttcttggatt tctaaaaagt 480
catcaatgac gtcattaatg cttttactgc tattcgcttt tgtacagccg tgtgcctcaa 540
tagtcgaaaa acgatgcggc ccaatcgata ttcgaaatag gccgtgggat attaagccgc 600
aatggtcgaa acttggtgat ccgaacgaaa aagatttggc tggtcagaga atggtcaact 660
gcacagtggt ggaaggttcg ctgacaatct catttgtact gaaacacaag acaaaagcac 720
aagaagaaat gcatcgaagt ctacagccaa gatattccca agacgaattt atcacttttc 780
cgcatctacg tgaaattact ggaactctgc tcgtttttga gactgaagga ttagtggatt 840
tgcgtaaaat tttcccaaat cttcgtgtaa ttggaggccg ttcgctgatt caacactatg 900
cgctgataat ttatcgaaat ccggatttgg agatcggtct tgacaagctt tccgtaattc 960
gaaatggtgg tgtacggata atcgataatc gaaaactgtg ctacacgaaa acgattgatt 1020
ggaaacattt gatcacttct tccatcaacg atgttgtcgt tgataatgct gccgagtacg 1080
ctgtcactga gactggattg atgtgcccac gtggagcttg cgaagaggat aaaggcgaat 1140
caaagtgtca ttatttggag gaaaagaatc aggaacaagg tgtcgaaaga gttcagagtt 1200
gttggtcgaa caccacttgc caaaagtctt gtgcttatga tcgtcttctt ccaacgaaag 1260
aaatcggacc gggatgtgat gcgaacggcg atcgatgtca cgatcaatgc gtgggcggtt 1320
gtgagcgtgt gaatgatgcc acagcatgcc acgcgtgcaa gaatgtctat cacaagggaa 1380
agtgtatcga aaagtgtgat gctcacctgt accttctcct tcaacgtcgt tgtgtgaccc 1440
gtgagcagtg tctgcagctg aatccggtgc tctcgaacaa aacagtgcct atcaaggcga 1500
```

```
cggcaggcct ttgctcggat aaatgtcccg atggttatca aatcaacccg gatgatcatc 1560
gagaatgccg aaaatgcgtt ggcaagtgtg agattgtgtg cgagatcaat cacgtcattg 1620
atacgtttcc gaaggcacag gcgatcaggc tatgcaatat tattgacgga aatctgacga 1680
tegagatteg eggaaaacag gattegggaa tegegteega gttgaaggat atatttgega 1740
acattcacac gatcaccggc tacctgttgg tacgtcaatc gtcaccgttt atctcgttga 1800
acatgttccg gaatttacga cgtattgagg caaagtcact gttcagaaat ctatatgcta 1860
tcacagtttt tgaaaatccg aatttaaaaa agctattcga ttcaacgacg gatttgacgc 1920
ttgatcgtgg aactgtgtca attgccaata acaagatgtt atgcttcaag tatatcaagc 1980
agctaatgtc aaagttaaat ataccactcg atccgataga tcaatcagaa gggacaaatg 2040
gtgagaaggn aatctgtgag gatatggcaa tcaacgtgag catcacagcg gtcaacgcgg 2100
actcggtctt ctttagttgg ccctcattca acattaccga tatagatcag cgaaagtttc 2160
teggetacga getettette aaagaagtee cacgaatega tgagaacatg acgategaag 2220
aggatcgaag tgcgtgtgtc gattcgtggc agagtgtctt caaacagtac tacgagacgt 2280
cgaacggtga accgaccccg gacattttta tggatattgg accgcgcgag cgaattcggc 2340
cgaatacgct ctacgcgtac tatgtggcga cgcagatggt gttgcatgcc ggtgcgaaga 2400
acggtgtatc gaagattggt tttgtgagga cgagctacta tacgcctgat cctccgacgt 2460
tggcactagc gcaagtcgat tcggacgcta ttcatattac gtgggaagcg ccgctccaac 2520
cgaacggaga cctcacgcat tacacaatta tgtggcgtga gaatgaagtg agcccgtacg 2580
aggaagccga aaagttttgt acagatgcaa gcacccccgc aaatcgacaa cgcacgaaag 2640
atccgaaaga gacgattgta gccgataagc cagtcgatat tccgtcatca cgtaccgtag 2700
ctccgacact tttgactatg atgggtcacg aagatcagca gaaaacgtgc gctgcaacgc 2760
ccggttgttg ttcgtgttcg gctatcgaag aatcatcgga acagaacaag aagaagcgac 2820
cggatccgat gtcggcgatc gaatcatctg catttgagaa taagctgttg gatgaggttt 2880
taatgccgag agacacgatg cgagtgagac gatcaattga agacgcgaat cgagtcagtg 2940
aagagttgga aaaagctgaa aatttgggaa aagctccaaa aactctcggt ggaaagaagc 3000
cgctgatcca tatttcgaag aagaagccgt cgagcagcag caccacatcc acaccggctc 3060
caacgatcgc atcaatgtat gccttaacaa ggaaaccgac tacggtgccg ggaacaagga 3120
ttcggctcta cgagatctac gaacctttac ccggaagctg ggcgattaat gtatcagctc 3180
tggcattgga taatagttat gtgatacgaa atttgaagca ttacacactt tatgcgattt 3240
ctctatccgc gtgccaaaac atgacagtac ccggagcatc ttgctcaata tcccatcgtg 3300
cgggagcatt gaaacgaaca aaacacatca cagacattga taaagtgttg aatgaaacaa 3360
ttgaatggag atttatgaat aatagtcaac aagtcaacgt gacgtgggat ccaccgactg 3420
aagtgaatgg tggaatattc ggttatgttg taaagcttaa gtcaaaagtc gatggatcaa 3480
ttgttatgac gagatgtgtc ggtgcgaaga gaggatattc aacacggaat cagggtgtcc 3540
tattccagaa tttggccgat ggacgttatt ttgtctcagt aacggcgacc tctgtacacg 3600
gegetggace ggaageegaa teeteegace caategtegt catgaegeea ggettettea 3660
ctgtggaaat cattctcggc atgcttctcg tctttttgat tttaatgtca attgccggtt 3720
gtataatcta ctactacatt caagtacgct acggcaaaaa agtgaaagct ctatctgact 3780
ttatgcaatt gaatcccgaa tattgtgtgg acaataagta caatgcagac gattgggagc 3840
tacggcagga tgatgttgtg ctcggacaac agtgtggaga gggatcattc ggaaaagtgt 3900
acctaggaac tggaaataat gttgtttctc tgatgggtga tcgtttcgga ccgtgtgcta 3960
 ttaagattaa tgtagatgat ccagcgtcga ctgagaatct caactatctc atggaagcta 4020
 atattatgaa gaactttaag actaacttta tcgtccaact gtacggagtt atctctactg 4080
 tacaaccagc gatggttgtg atggaaatga tggatcttgg aaatctccgt gactatctcc 4140
 gatcgaaacg cgaagacgaa gtgttcaatg agacggactg caactttttc gacataatcc 4200
 cgagggataa attccatgag tgggccgcac agatttgtga tggtatggcg tacctggagt 4260
 cgctcaagtt ttgccatcga gatctcgccg cacgtaattg catgataaat cgggatgaga 4320
 ctgtcaagat tggagatttc ggaatggctc gtgatctatt ctatcatgac tattataagc 4380
 catcgggcaa gcgtatgatg cctgttcgat ggatgtcacc cgagtcgttg aaagacggaa 4440
 agtttgactc gaaatctgat gtttggagct tcggagttgt tctctatgaa atggttacac 4500
 teggtgetea gecatatatt ggtttgagta atgatgaggt gttgaattat attggaatgg 4560
 cccggaaggt tatcaagaag cccgaatgtt gtgaaaacta ttggtataag gtgatgaaaa 4620
 tgtgctggag atactcacct cgggatcgtc cgacgttcct ccagctcgtt catcttctag 4680
 cagctgaagc ttcaccagaa ttccgagatt tatcatttgt cctaaccgat aatcaaatga 4740
 teettgaega tteagaagea etggatettg atgatattga tgataetgat atgaatgate 4800
 aggttgtcga ggtggcaccg gatgttgaga acgtcgaggt tcagagtgat tcggaacgtc 4860
 ggaatacgga ttcaataccg ttgaaacagt ttaagacgat ccctccgatc aatgcgacga 4920
 cgagtcattc gacaatatcg attgatgaga caccgatgaa agcgaagcag cgagaaggat 4980
```

```
cgctggatga ggagtacgca ttgatgaatc atagtggagg tccgagtgat gcggaagttc 5040 ggacgtatgc tggtgatgga gattatgtgg agagagatgt tcgagagaat gatgtgccaa 5100 cgcgacgaaa tactggtgca tcaacatcaa gttacacagg tggtggtcca tattgcctaa 5160 caaatcgtgg tggttcaaat gaacgaggag ccggtttcgg tgaagcagta cgattaactg 5220 atggtgtgg aagtggacat ttaaatgatg atgattatgt tgaaaaagag atatcatcca 5280 tggatacgcg ccggagcacg ggcgcctcga gctcttccta cggtgttcca cagacgaatt 5340 ggagtggaaa tcgtggtgcc acgtattata cgagtaaagc tcaacaggca gcaactgcag 5400 cagcagcagc agcagcagc ctccaacagc aacaaaatgg tggtcgaggc gatcgattaa 5460 ctcaactacc cggaactgga catttacaat cgacacgtgg tggacaagat ggagattata 5520 ttgaaactga accgaaaat tatagaaata atggatctcc atcgcgaaac ggcgacagcc 5580 gtgacatttt caacggacgt tcggctttcg gtgaaaatga gcatctaatc gaggataatg 5640 agcatcatcc acttgtctga aacccccaaa aaatcccgcc tcttaaatta taaattatct 5700 cccacattat catactcta cacgaatatc tctgtattt tcttgtatt tcttgtat tacccc 5816
```

<210> 12 <211> 1724 <212> PRT <213> Caenorhabditis elegans

<400> 12

10 Ala Ser Ile Val Glu Lys Arg Cys Gly Pro Ile Asp Ile Arg Asn Arg 20 25 Pro Trp Asp Ile Lys Pro Gln Trp Ser Lys Leu Gly Asp Pro Asn Glu 40 45 Lys Asp Leu Ala Gly Gln Arg Met Val Asn Cys Thr Val Val Glu Gly 55 60 Ser Leu Thr Ile Ser Phe Val Leu Lys His Lys Thr Lys Ala Gln Glu 70 75 Glu Met His Arg Ser Leu Gln Pro Arg Tyr Ser Gln Asp Glu Phe Ile 90 85 Thr Phe Pro His Leu Arg Glu Ile Thr Gly Thr Leu Leu Val Phe Glu 100 105 110 Thr Glu Gly Leu Val Asp Leu Arg Lys Ile Phe Pro Asn Leu Arg Val 120 Ile Gly Gly Arg Ser Leu Ile Gln His Tyr Ala Leu Ile Ile Tyr Arg 135 140 Asn Pro Asp Leu Glu Ile Gly Leu Asp Lys Leu Ser Val Ile Arg Asn 150 155 Gly Gly Val Arg Ile Ile Asp Asn Arg Lys Leu Cys Tyr Thr Lys Thr 170 165 175 Ile Asp Trp Lys His Leu Ile Thr Ser Ser Ile Asn Asp Val Val Val 180 185 190 Asp Asn Ala Ala Glu Tyr Ala Val Thr Glu Thr Gly Leu Met Cys Pro 200 205 Arg Gly Ala Cys Glu Glu Asp Lys Gly Glu Ser Lys Cys His Tyr Leu 215 220 Glu Glu Lys Asn Gln Glu Gln Gly Val Glu Arg Val Gln Ser Cys Trp 230 235 Ser Asn Thr Thr Cys Gln Lys Ser Cys Ala Tyr Asp Arg Leu Leu Pro 245 250 255 Thr Lys Glu Ile Gly Pro Gly Cys Asp Ala Asn Gly Asp Arg Cys His 265 270 Asp Gln Cys Val Gly Gly Cys Glu Arg Val Asn Asp Ala Thr Ala Cys

Met Thr Ser Leu Met Leu Leu Leu Phe Ala Phe Val Gln Pro Cys

		275					280					285			
His	Ala 290	Cys	Lys	Asn	Val	Tyr 295	His	Lys	Gly	Lys	Cys 300	Ile	Glu	Lys	Cys
Asp 305	Ala	His	Leu	Tyr	Leu 310	Leu	Leu	Gln	Arg	Arg 315	Cys	Val	Thr	Arg	Glu 320
Gln	Cys	Leu	Gln	Leu 325	Asn	Pro	Val	Leu	Ser 330	Asn	Lys	Thr	Val	Pro 335	Ile
Lys	Ala	Thr	Ala 340	Gly	Leu	Cys	Ser	Asp 345	Lys	Cys	Pro	Asp	Gly 350	Tyr	Gln
Ile	Asn	Pro 355	Asp	Asp	His	Arg	Glu 360	Cys	Arg	Lys	Cys	Val 365	Gly	Lys	Cys
	Ile 370					375					380				
385	Ala				390					395					400
	Arg			405					410					415	
	Ala		420					425					430		
	Pro	435					440		_			445	_		
	Lys 450					455					460				
465	Asn			-	470					475	_				480
•	Gly			485					490			-		495	_
	Lys		500					505					510		
	Ser	Glu 515	Gly	Thr	Asn	GLY	G1u 520	Lys	Ala	Ile	Cys	Glu 525	Asp	Met	Ala
								_		_			1		
	Asn 530					535					540				
Trp 545	530 Pro	Ser	Phe	Asn	Ile 550	535 Thr	Asp	Ile	Asp	Gln 555	540 Arg	Lys	Phe	Leu	Gly 560
Trp 545 Tyr	530 Pro Glu	Ser Leu	Phe Phe	Asn Phe 565	Ile 550 Lys	535 Thr Glu	Asp Val	Ile Pro	Asp Arg 570	Gln 555 Ile	540 Arg Asp	Lys Glu	Phe Asn	Leu Met 575	Gly 560 Thr
Trp 545 Tyr Ile	530 Pro Glu Glu	Ser Leu Glu	Phe Phe Asp 580	Asn Phe 565 Arg	Ile 550 Lys Ser	535 Thr Glu Ala	Asp Val Cys	Ile Pro Val 585	Asp Arg 570 Asp	Gln 555 Ile Ser	540 Arg Asp Trp	Lys Glu Gln	Phe Asn Ser 590	Leu Met 575 Val	Gly 560 Thr
Trp 545 Tyr Ile Lys	530 Pro Glu Glu Gln	Ser Leu Glu Tyr 595	Phe Phe Asp 580 Tyr	Asn Phe 565 Arg Glu	Ile 550 Lys Ser	535 Thr Glu Ala Ser	Asp Val Cys Asn 600	Ile Pro Val 585 Gly	Asp Arg 570 Asp Glu	Gln 555 Ile Ser Pro	540 Arg Asp Trp	Lys Glu Gln Pro 605	Phe Asn Ser 590 Asp	Leu Met 575 Val	Gly 560 Thr Phe
Trp 545 Tyr Ile Lys Met	530 Pro Glu Glu Gln Asp 610	Ser Leu Glu Tyr 595 Ile	Phe Phe Asp 580 Tyr	Asn Phe 565 Arg Glu Pro	Ile 550 Lys Ser Thr	535 Thr Glu Ala Ser Glu 615	Asp Val Cys Asn 600 Arg	Ile Pro Val 585 Gly Ile	Asp 570 Asp Glu Arg	Gln 555 Ile Ser Pro	540 Arg Asp Trp Thr Asn 620	Lys Glu Gln Pro 605 Thr	Phe Asn Ser 590 Asp	Leu Met 575 Val Ile Tyr	Gly 560 Thr Phe Phe
Trp 545 Tyr Ile Lys Met Tyr 625	530 Pro Glu Glu Gln Asp 610 Tyr	Ser Leu Glu Tyr 595 Ile Val	Phe Phe Asp 580 Tyr Gly Ala	Asn Phe 565 Arg Glu Pro Thr	Ile 550 Lys Ser Thr Arg Gln 630	535 Thr Glu Ala Ser Glu 615 Met	Asp Val Cys Asn 600 Arg Val	Ile Pro Val 585 Gly Ile Leu	Asp 570 Asp Glu Arg	Gln 555 Ile Ser Pro Pro Ala 635	540 Arg Asp Trp Thr Asn 620 Gly	Lys Glu Gln Pro 605 Thr	Phe Asn Ser 590 Asp Leu Lys	Leu Met 575 Val Ile Tyr Asn	Gly 560 Thr Phe Phe Ala Gly 640
Trp 545 Tyr Ile Lys Met Tyr 625 Val	530 Pro Glu Glu Gln Asp 610 Tyr	Ser Leu Glu Tyr 595 Ile Val Lys	Phe Phe Asp 580 Tyr Gly Ala Ile	Asn Phe 565 Arg Glu Pro Thr Gly 645	Ile 550 Lys Ser Thr Arg Gln 630 Phe	535 Thr Glu Ala Ser Glu 615 Met Val	Asp Val Cys Asn 600 Arg Val	Ile Pro Val 585 Gly Ile Leu Thr	Asp 570 Asp Glu Arg His Ser 650	Gln 555 Ile Ser Pro Pro Ala 635 Tyr	540 Arg Asp Trp Thr Asn 620 Gly	Lys Glu Gln Pro 605 Thr Ala Thr	Phe Asn Ser 590 Asp Leu Lys Pro	Leu Met 575 Val Ile Tyr Asn Asp 655	Gly 560 Thr Phe Phe Ala Gly 640 Pro
Trp 545 Tyr Ile Lys Met Tyr 625 Val	530 Pro Glu Glu Gln Asp 610 Tyr Ser	Ser Leu Glu Tyr 595 Ile Val Lys Leu	Phe Phe Asp 580 Tyr Gly Ala Ile Ala 660	Asn Phe 565 Arg Glu Pro Thr Gly 645 Leu	Ile 550 Lys Ser Thr Arg Gln 630 Phe	535 Thr Glu Ala Ser Glu 615 Met Val Gln	Asp Val Cys Asn 600 Arg Val Arg	Ile Pro Val 585 Gly Ile Leu Thr Asp 665	Asp 570 Asp Glu Arg His Ser 650 Ser	Gln 555 Ile Ser Pro Pro Ala 635 Tyr	540 Arg Asp Trp Thr Asn 620 Gly Tyr	Lys Glu Gln Pro 605 Thr Ala Thr	Phe Asn Ser 590 Asp Leu Lys Pro His 670	Leu Met 575 Val Ile Tyr Asn Asp 655 Ile	Gly 560 Thr Phe Phe Ala Gly 640 Pro
Trp 545 Tyr Ile Lys Met Tyr 625 Val Pro	530 Pro Glu Glu Gln Asp 610 Tyr Ser Thr Glu	Ser Leu Glu Tyr 595 Ile Val Lys Leu Ala 675	Phe Phe Asp 580 Tyr Gly Ala Ile Ala 660 Pro	Asn Phe 565 Arg Glu Pro Thr Gly 645 Leu Leu	Ile 550 Lys Ser Thr Arg Gln 630 Phe Ala Gln	535 Thr Glu Ala Ser Glu 615 Met Val Gln Pro	Asp Val Cys Asn 600 Arg Val Arg Val Asn 680	Ile Pro Val 585 Gly Ile Leu Thr Asp 665 Gly	Asp Arg 570 Asp Glu Arg His Ser 650 Ser Asp	Gln 555 Ile Ser Pro Pro Ala 635 Tyr Asp Leu	540 Arg Asp Trp Thr Asn 620 Gly Tyr Ala Thr	Lys Glu Gln Pro 605 Thr Ala Thr Ile His 685	Phe Asn Ser 590 Asp Leu Lys Pro His 670 Tyr	Leu Met 575 Val Ile Tyr Asn Asp 655 Ile Thr	Gly 560 Thr Phe Phe Ala Gly 640 Pro Thr
Trp 545 Tyr Ile Lys Met Tyr 625 Val Pro Trp Met	S30 Pro Glu Glu Gln Asp 610 Tyr Ser Thr Glu Trp 690	Ser Leu Glu Tyr 595 Ile Val Lys Leu Ala 675 Arg	Phe Phe Asp 580 Tyr Gly Ala Ile Ala 660 Pro Glu	Asn Phe 565 Arg Glu Pro Thr Gly 645 Leu Leu Asn	Ile 550 Lys Ser Thr Arg Gln 630 Phe Ala Gln Glu	535 Thr Glu Ala Ser Glu 615 Met Val Gln Pro Val 695	Asp Val Cys Asn 600 Arg Val Arg Val Asn 680 Ser	Ile Pro Val 585 Gly Ile Leu Thr Asp 665 Gly Pro	Asp Arg 570 Asp Glu Arg His Ser 650 Ser Asp	Gln 555 Ile Ser Pro Pro Ala 635 Tyr Asp Leu Glu	540 Arg Asp Trp Thr Asn 620 Gly Tyr Ala Thr Glu 700	Lys Glu Gln Pro 605 Thr Ala Thr Ile His 685 Ala	Phe Asn Ser 590 Asp Leu Lys Pro His 670 Tyr Glu	Leu Met 575 Val Ile Tyr Asn Asp 655 Ile Thr	Gly 560 Thr Phe Phe Ala Gly 640 Pro Thr Ile
Trp 545 Tyr Ile Lys Met Tyr 625 Val Pro Trp Met Cys 705	S30 Pro Glu Glu Gln Asp 610 Tyr Ser Thr Glu Trp 690 Thr	Ser Leu Glu Tyr 595 Ile Val Lys Leu Ala 675 Arg Asp	Phe Phe Asp 580 Tyr Gly Ala Ile A660 Pro Glu Ala	Asn Phe 565 Arg Glu Pro Thr Gly 645 Leu Leu Asn Ser	Ile 550 Lys Ser Thr Arg Gln 630 Phe Ala Gln Glu Thr 710	535 Thr Glu Ala Ser Glu 615 Met Val Gln Pro Val 695 Pro	Asp Val Cys Asn 600 Arg Val Arg Val Asn 680 Ser Ala	Ile Pro Val 585 Gly Ile Leu Thr Asp 665 Gly Pro Asn	Asp Arg 570 Asp Glu Arg His Ser 650 Ser Asp Tyr Arg	Gln 555 Ile Ser Pro Pro Ala 635 Tyr Asp Leu Glu Gln 715	540 Arg Asp Trp Thr Asn 620 Gly Tyr Ala Thr Glu 700 Arg	Lys Glu Gln Pro 605 Thr Ala Thr Ile His 685 Ala Thr	Phe Asn Ser 590 Asp Leu Lys Pro His 670 Tyr Glu Lys	Leu Met 575 Val Ile Tyr Asn Asp 655 Ile Thr Lys Asp	Gly 560 Thr Phe Phe Ala Gly 640 Pro Thr Ile Phe Pro 720
Trp 545 Tyr Ile Lys Met Tyr 625 Val Pro Trp Met Cys 705 Lys	S30 Pro Glu Glu Gln Asp 610 Tyr Ser Thr Glu Trp 690	Ser Leu Glu Tyr 595 Ile Val Lys Leu Ala 675 Arg Asp Thr	Phe Phe Asp 580 Tyr Gly Ala Ile A660 Pro Glu Ala Ile	Asn Phe 565 Arg Glu Pro Thr Gly 645 Leu Asn Ser Val 725	Ile 550 Lys Ser Thr Arg Gln 630 Phe Ala Gln Glu Thr 710 Ala	S35 Thr Glu Ala Ser Glu 615 Met Val Gln Pro Val 695 Pro Asp	Asp Val Cys Asn 600 Arg Val Arg Val Asn 680 Ser Ala Lys	Ile Pro Val 585 Gly Ile Leu Thr Asp 665 Gly Pro Asn Pro	Asp Arg 570 Asp Glu Arg His Ser 650 Ser Asp Tyr Arg Val 730	Gln 555 Ile Ser Pro Pro Ala 635 Tyr Asp Leu Glu Gln 715 Asp	Asp Trp Thr Asn 620 Gly Tyr Ala Thr Glu 700 Arg	Lys Glu Gln Pro 605 Thr Ala Thr Ile His 685 Ala Thr	Phe Asn Ser 590 Asp Leu Lys Pro His 670 Tyr Glu Lys Ser	Leu Met 575 Val Ile Tyr Asn Asp 655 Ile Thr Lys Asp Ser 735	Gly 560 Thr Phe Phe Ala Gly 640 Pro Thr Ile Phe Pro 720 Arg

Lys Thr Cys Ala Ala Thr Pro Gly Cys Cys Ser Cys Ser Ala Ile Glu Glu Ser Ser Glu Gln Asn Lys Lys Arg Pro Asp Pro Met Ser Ala Ile Glu Ser Ser Ala Phe Glu Asn Lys Leu Leu Asp Glu Val Leu Met Pro Arg Asp Thr Met Arg Val Arg Arg Ser Ile Glu Asp Ala Asn Arg Val Ser Glu Glu Leu Glu Lys Ala Glu Asn Leu Gly Lys Ala Pro Lys Thr Leu Gly Gly Lys Lys Pro Leu Ile His Ile Ser Lys Lys Pro Ser Ser Ser Ser Thr Thr Ser Thr Pro Ala Pro Thr Ile Ala Ser Met Tyr Ala Leu Thr Arg Lys Pro Thr Thr Val Pro Gly Thr Arg Ile Arg Leu Tyr Glu Ile Tyr Glu Pro Leu Pro Gly Ser Trp Ala Ile Asn Val Ser Ala Leu Ala Leu Asp Asn Ser Tyr Val Ile Arg Asn Leu Lys His Tyr Thr Leu Tyr Ala Ile Ser Leu Ser Ala Cys Gln Asn Met Thr Val Pro Gly Ala Ser Cys Ser Ile Ser His Arg Ala Gly Ala Leu Lys Arg Thr Lys His Ile Thr Asp Ile Asp Lys Val Leu Asn Glu Thr Ile Glu Trp Arg Phe Met Asn Asn Ser Gln Gln Val Asn Val Thr Trp Asp Pro Pro Thr Glu Val Asn Gly Gly Ile Phe Gly Tyr Val Val Lys Leu Lys Ser Lys Val Asp Gly Ser Ile Val Met Thr Arg Cys Val Gly Ala Lys Arg Gly Tyr Ser Thr Arg Asn Gln Gly Val Leu Phe Gln Asn Leu Ala 1010 1015 Asp Gly Arg Tyr Phe Val Ser Val Thr Ala Thr Ser Val His Gly Ala 1030 ,1035 Gly Pro Glu Ala Glu Ser Ser Asp Pro Ile Val Val Met Thr Pro Gly 1050 1055 Phe Phe Thr Val Glu Ile Ile Leu Gly Met Leu Leu Val Phe Leu Ile Leu Met Ser Ile Ala Gly Cys Ile Ile Tyr Tyr Tyr Ile Gln Val Arg 1080 1085 Tyr Gly Lys Lys Val Lys Ala Leu Ser Asp Phe Met Gln Leu Asn Pro Glu Tyr Cys Val Asp Asn Lys Tyr Asn Ala Asp Asp Trp Glu Leu Arg Gln Asp Asp Val Val Leu Gly Gln Gln Cys Gly Glu Gly Ser Phe Gly Lys Val Tyr Leu Gly Thr Gly Asn Asn Val Val Ser Leu Met Gly Asp Arg Phe Gly Pro Cys Ala Ile Lys Ile Asn Val Asp Asp Pro Ala Ser 1155 1160 Thr Glu Asn Leu Asn Tyr Leu Met Glu Ala Asn Ile Met Lys Asn Phe Lys Thr Asn Phe Ile Val Gln Leu Tyr Gly Val Ile Ser Thr Val Gln Pro Ala Met Val Val Met Glu Met Met Asp Leu Gly Asn Leu Arg Asp

				1205	5				1210)				1215	ō
Tyr	Leu	Arg	Ser 1220		Arg	Glu	Asp	Glu 1225		Phe	Asn	Glu	Thr 1230		Cys
Asn	Phe	Phe 123!	qaA o	Ile	Ile	Pro	Arg 1240		Lys	Phe	His	Glu 1245		Ala	Ala
Gln	Ile 1250		Asp	Gly	Met	Ala 1255	_	Leu	Glu	Ser	Leu 126		Phe	Cys	His
Arg 1265	_	Leu	Ala	Ala	Arg 1270		Cys	Met	Ile	Asn 1275		Asp	Glu	Thr	Val 1280
Lys	Ile	Gly	Asp	Phe 1285		Met	Ala	Arg	Asp 1290		Phe	Tyr	His	Asp 1295	_
Tyr	Lys	Pro	Ser 1300		Lys	Arg	Met	Met 1305		Val	Arg	Trp	Met 131		Pro
Glu	Ser	Leu 131	Lys S	Asp	Gly	Lys	Phe 1320		Ser	Lys	Ser	Asp 1325		Trp	Ser
Phe	Gly 1330		Val	Leu	Tyr	Glu 1335		Val	Thr	Leu	Gly 1340		Gln	Pro	Tyr
1345	5		Ser		1350)				1355	5				1360
Lys	Val	Ile	Lys	Lys 1365		Glu	Cys	Cys	Glu 1370		Tyr	Trp	Tyr	Lys 1375	
Met	Lys	Met	Cys 1380		Arg	Tyr	Ser	Pro 1385		Asp	Arg	Pro	Thr 1390		Leu
		1395					1400)				1405	ō		
	1410)	Val			1415	5				1420)	_		
1425	5		Leu		1430)				1435	5				1440
			Ala	1445	5				1450)				1455	5
			Asn 1460)				1465	5				1470)	
		1475					1480)				1485	5		
	1490)	Lys			1495	5				1500)			
1505	5		Asn		1510)				1515	5				1520
			Asp	1525	5				1530)				1535	5
			Arg 1540)				1545	5				1550)	
		1555					1560)				1565	5		
	1570)	Gly			1575	5				1580)			
1585	5		Asp		1590)				1595	5				1600
			Ser	1605	5				1610)				1615	5
			Ser 1620)				1625	5				1630)	
		1635					1640)				1645	5		
	1650)	Gly			1655	5				1660)		_	
Gly	His	Leu	Gln	Ser	Thr	Arg	Gly	GLy	Gln	Asp	GLy	Asp	Tyr	Ile	Glu

```
1675
                   1670
1665
Thr Glu Pro Lys Asn Tyr Arg Asn Asn Gly Ser Pro Ser Arg Asn Gly
           1685
                            1690
Asn Ser Arg Asp Ile Phe Asn Gly Arg Ser Ala Phe Gly Glu Asn Glu
           1700
                              1705
His Leu Ile Glu Asp Asn Glu His His Pro Leu Val
                            1720
<210> 13
<211> 139
<212> PRT
<213> Caenorhabditis elegans
<400> 13
Thr Ser Gly Ser Gly Met Gly Pro Thr Thr Leu His Lys Leu Thr Ile
                                    10
Gly Gly Gln Ile Arg Leu Thr Gly Arg Val Gly Ser Gly Arg Phe Gly
                                25
Asn Val Ser Arg Gly Asp Tyr Arg Gly Glu Ala Val Ala Val Lys Val
                                                45
Phe Asn Ala Leu Asp Glu Pro Ala Phe His Lys Glu Thr Glu Ile Phe
                                            60
                        55
Glu Thr Arg Met Leu Arg His Pro Asn Val Leu Arg Tyr Ile Gly Ser
                                        75
                    70
Asp Arg Val Asp Thr Gly Phe Val Thr Glu Leu Trp Leu Val Thr Glu
                                    90
Tyr His Pro Ser Gly Ser Leu His Asp Phe Leu Leu Glu Asn Thr Val
                                105
            100
Asn Ile Glu Thr Tyr Tyr Asn Leu Met Arg Ser Thr Ala Ser Gly Leu
                            120
Ala Phe Leu His Asn Gln Ile Gly Gly Ser Lys
                        135
    130
<210> 14
<211> 62
<212> PRT
<213> Caenorhabditis elegans
<400> 14
Glu Asp Ala Ala Ser Asp Ile Ile Ala Asn Glu Asn Tyr Lys Cys Gly
                                     10
Thr Val Arg Tyr Leu Ala Pro Glu Ile Leu Asn Ser Thr Met Gln Phe
                                                     30
            20
 Thr Val Phe Glu Ser Tyr Gln Cys Ala Asp Val Tyr Ser Phe Ser Leu
                             40
 Val Met Trp Glu Thr Leu Cys Arg Cys Glu Asp Gly Asp Val
 <210> 15
```

<211> 31

<212> PRT

<213> Caenorhabditis elegans

Lys Pro Ala Met Ala His Arg Asp Ile Lys Ser Lys Asn Ile Met Val

```
15
                                    10
Lys Asn Asp Leu Thr Cys Ala Ile Gly Asp Leu Gly Leu Ser Leu
                                25
            20
<210> 16
<211> 72
<212> PRT
<213> Caenorhabditis elegans
<400> 16
Ile Pro Tyr Ile Glu Trp Thr Asp Arg Asp Pro Gln Asp Ala Gln Met
Phe Asp Val Val Cys Thr Arg Arg Leu Arg Pro Thr Glu Asn Pro Leu
                                25
Trp Lys Asp His Pro Glu Met Lys His Ile Met Glu Ile Ile Lys Thr
                            40
Cys Trp Asn Gly Asn Pro Ser Ala Arg Phe Thr Ser Tyr Ile Cys Arg
                        55
Lys Arg Met Asp Glu Arg Gln Gln
<210> 17
<211> 150
<212> PRT
<213> Caenorhabditis elegans
<400> 17
Tyr Phe Glu Ser Val Asp Arg Phe Leu Tyr Ser Cys Val Gly Tyr Ser
 1
Val Ala Thr Tyr Ile Met Gly Ile Lys Asp Arg His Ser Asp Asn Leu
                                 25
             20
Met Leu Thr Glu Asp Gly Lys Tyr Val His Ile Asp Phe Gly His Ile
                             40
Leu Gly His Gly Lys Thr Lys Leu Gly Ile Gln Arg Asp Arg Gln Pro
                                             60
                         55
 Phe Ile Leu Thr Glu His Phe Met Thr Val Ile Arg Ser Gly Lys Ser
                                         75
                     70
Val Asp Gly Asn Ser His Glu Leu Gln Lys Phe Lys Thr Leu Cys Val
                                                         95
                                     90
 Glu Ala Tyr Glu Val Met Trp Asn Asn Arg Asp Leu Phe Val Ser Leu
                                 105
             100
 Phe Thr Leu Met Leu Gly Met Glu Leu Pro Glu Leu Ser Thr Lys Ala
                                                 125
                             120
         115
 Asp Leu Asp His Leu Lys Lys Thr Leu Phe Cys Asn Gly Glu Ser Lys
                         135
 Glu Glu Ala Arg Lys Phe
                     150
 <210> 18
 <211> 113
 <212> PRT
 <213> Caenorhabditis elegans
 Ser Pro Leu Asp Pro Val Tyr Lys Leu Gly Glu Met Ile Ile Asp Lys
```

<210> 19 <211> 106 <212> PRT <213> Caenorhabditis elegans

<400> 19 Leu Ala Phe Val Trp Thr Asp Arg Glu Asn Phe Ser Glu Leu Tyr Val 10 Met Leu Glu Lys Trp Lys Pro Pro Ser Val Ala Ala Ala Leu Thr Leu 25 Leu Gly Lys Arg Cys Thr Asp Arg Val Ile Arg Lys Phe Ala Val Glu 40 Lys Leu Asn Glu Gln Leu Ser Pro Val Thr Phe His Leu Phe Ile Leu 55 Pro Leu Ile Gln Ala Leu Lys Tyr Glu Pro Arg Ala Gln Ser Glu Val 75 70 Gly Met Met Leu Leu Thr Arg Ala Leu Cys Asp Tyr Arg Ile Gly His 90 85 Arg Leu Phe Trp Leu Leu Arg Ala Glu Ile 100

<210> 20 <211> 139 <212> PRT <213> Caenorhabditis elegans

<400> 20 Glu Tyr Trp Ile Val Thr Glu Phe His Glu Arg Leu Ser Leu Tyr Glu 10 Leu Leu Lys Asn Asn Val Ile Ser Ile Thr Ser Ala Asn Arg Ile Ile 25 Met Ser Met Ile Asp Gly Leu Gln Phe Leu His Asp Asp Arg Pro Tyr 40 Phe Phe Gly His Pro Lys Lys Pro Ile Ile His Arg Asp Ile Lys Ser 55 60 Lys Asn Ile Leu Val Lys Ser Asp Met Thr Thr Cys Ile Ala Asp Phe 75 70 Gly Leu Ala Arg Ile Tyr Ser Tyr Asp Ile Glu Gln Ser Asp Leu Leu 90 Gly Gln Val Gly Thr Lys Arg Tyr Met Ser Pro Glu Met Leu Glu Gly

```
100
                               105
Ala Thr Glu Phe Thr Pro Thr Ala Phe Lys Ala Met Asp Val Tyr Ser
                120
Met Gly Leu Val Met Trp Glu Val Ile Ser Arg
                      135
  130
<210> 21
<211> 61
<212> PRT
<213> Caenorhabditis elegans
<400> 21
Ile Gly Phe Asp Pro Thr Ile Gly Arg Met Arg Asn Tyr Val Val Ser
                                    10
Lys Lys Glu Arg Pro Gln Trp Arg Asp Glu Ile Ile Lys His Glu Tyr
                                25
Met Ser Leu Leu Lys Lys Val Thr Glu Glu Met Trp Asp Pro Glu Ala
                            40
Cys Ala Arg Ile Thr Ala Gly Cys Ala Phe Ala Arg Val
                        55
<210> 22
<211> 20
<212> PRT
<213> Caenorhabditis elegans
<400> 22
Pro Ile Thr Asp Phe Gln Leu Ile Ser Lys Gly Arg Phe Gly Lys Val
Phe Lys Ala Gln
           2.0
<210> 23
<211> 163
<212> PRT
<213> Caenorhabditis elegans
<400> 23
Thr Asp Ser Glu Thr Arg Ser Arg Phe Ser Leu Gly Trp Tyr Asn Asn
                                    10
Pro Asn Arg Ser Pro Gln Thr Ala Glu Val Arg Gly Leu Ile Gly Lys
           20
                                25
Gly Val Arg Phe Tyr Leu Leu Ala Gly Glu Val Tyr Val Glu Asn Leu
                            40
Cys Asn Ile Pro Val Phe Val Gln Ser Ile Gly Ala Asn Met Lys Asn
                       55
                                            60
Gly Phe Gln Leu Asn Thr Val Ser Lys Leu Pro Pro Thr Gly Thr Met
                   70
Lys Val Phe Asp Met Arg Leu Phe Ser Lys Gln Leu Arg Thr Ala Ala
               85
Glu Lys Thr Tyr Gln Asp Val Tyr Cys Leu Ser Arg Met Cys Thr Val
           100
                               105
Arg Val Ser Phe Cys Lys Gly Trp Gly Glu His Tyr Arg Arg Ser Thr
                            120
Val Leu Arg Ser Pro Val Trp Phe Gln Ala His Leu Asn Asn Pro Met
```

```
130
                        135
His Trp Val Asp Ser Val Leu Thr Cys Met Gly Ala Pro Pro Arg Ile
                    150
                                    155
Cys Ser Ser
<210> 24
<211> 44
<212> PRT
<213> Caenorhabditis elegans
<400> 24
Arg Ala Phe Arg Phe Pro Val Ile Arg Tyr Glu Ser Gln Val Lys Ser
                5
                                    10
Ile Leu Thr Cys Arg His Ala Phe Asn Ser His Ser Arg Asn Val Cys
        20
                                25
Leu Asn Pro Tyr His Tyr Arg Trp Val Glu Leu Pro
<210> 25
<211> 38
<212> PRT
<213> Caenorhabditis elegans
<400> 25
Val Glu Tyr Glu Glu Ser Pro Ser Trp Leu Lys Leu Ile Tyr Tyr Glu
Glu Gly Thr Met Ile Gly Glu Lys Ala Asp Val Glu Gly His His Cys
                                25
Leu Ile Asp Gly Phe Thr
        35
<210> 26
<211> 60
<212> PRT
<213> Caenorhabditis elegans
<400> 26
Asn Leu Ala Glu Thr Gly His Ser Lys Ile Met Arg Ala Ala His Lys
Val Ser Asn Pro Glu Ile Gly Tyr Cys Cys His Pro Thr Glu Tyr Asp
                                25
Tyr Ile Lys Leu Ile Tyr Val Asn Arg Asp Gly Arg Val Ser Ile Ala
                            40
Asn Val Asn Gly Met Ile Ala Lys Lys Cys Gly Cys
  50
                        55
<210> 27
<211> 20
<212> PRT
<213> Caenorhabditis elegans
<400> 27
```

Asp Trp Ile Val Ala Pro Pro Arg Tyr Asn Ala Tyr Met Cys Arg Gly

```
15
                                   10
Asp Cys His Tyr
            20
<210> 28
<211> 43
<212> PRT
<213> Caenorhabditis elegans
<400> 28
Val Cys Asn Ala Glu Ala Gln Ser Lys Gly Cys Cys Leu Tyr Asp Leu
                                   10
                5
Glu Ile Glu Phe Glu Lys Ile Gly Trp Asp Trp Ile Val Ala Pro Pro
                                25
            20
Arg Tyr Asn Ala Tyr Met Cys Arg Gly Asp Cys
        35
<210> 29
<211> 70
<212> PRT
<213> Caenorhabditis elegans
<400> 29
Asp Cys His Tyr Asn Ala His His Phe Asn Leu Ala Glu Thr Gly His
                                    10
                 5
1
Ser Lys Ile Met Arg Ala Ala His Lys Val Ser Asn Pro Glu Ile Gly
                                25
            20
Tyr Cys Cys His Pro Thr Glu Tyr Asp Tyr Ile Lys Leu Ile Tyr Val
                                                45
                            40
        35
Asn Arg Asp Gly Arg Val Ser Ile Ala Asn Val Asn Gly Met Ile Ala
                         55
Lys Lys Cys Gly Cys Ser
65
<210> 30
<211> 35
<212> PRT
<213> Caenorhabditis elegans
<400> 30
Cys Cys Leu Tyr Asp Leu Glu Ile Glu Phe Glu Lys Ile Gly Trp Asp
                                     10
 1
Trp Ile Val Ala Pro Pro Arg Tyr Asn Ala Tyr Met Cys Arg Gly Asp
                                 25
             20
 Cys His Tyr
        35
 <210> 31
 <211> 23
 <212> DNA
 <213> Artificial Sequence
 <220>
 <223> Degenerate probe
```

35

```
<221> misc_feature
<222> (1)...(23)
<223> n = A,T,C or G
<400> 31
                                                                   23
ggntgggayt rnrtnrtngc ncc
<210> 32
<211> 18
<212> DNA
<213> Artificial Sequence
<220>
<223> Degenerate probe
<221> misc_feature
<222> (1)...(18)
<223> n = A,T,C or G
<400> 32
                                                                   18
tgytgynnnc cnacngar
<210> 33
<211> 127
<212> PRT
<213> Caenorhabditis elegans
<400> 33
Lys Phe His Glu Trp Ala Ala Gln Ile Cys Asp Gly Met Ala Tyr Leu
1
                 5
                                    10
Glu Ser Leu Lys Phe Cys His Arg Asp Leu Ala Ala Arg Asn Cys Met
           20
                                25
Ile Asn Arg Asp Glu Thr Val Lys Ile Gly Asp Phe Gly Met Ala Arg
                            40
                                                 45
Asp Leu Phe Tyr His Asp Tyr Tyr Lys Pro Ser Gly Lys Arg Met Met
                        55
                                            60
Pro Val Arg Trp Met Ser Pro Glu Ser Leu Lys Asp Gly Lys Phe Asp
                                        75
                    70
Ser Lys Ser Asp Val Trp Ser Phe Gly Val Val Leu Tyr Glu Met Val
                85
                                    90
Thr Leu Gly Ala Gln Pro Tyr Ile Gly Leu Ser Asn Asp Glu Val Leu
            100
                                105
                                                     110
Asn Tyr Ile Gly Met Ala Arg Lys Val Ile Lys Lys Pro Glu Cys
<210> 34
<211> 131
<212> PRT
<213> Caenorhabditis elegans
<400> 34
Asn Thr Thr Cys Gln Lys Ser Cys Ala Tyr Asp Arg Leu Leu Pro Thr
Lys Glu Ile Gly Pro Gly Cys Asp Ala Asn Gly Asp Arg Cys His Asp
            20
                                25
                                                     30
Gln Cys Val Gly Gly Cys Glu Arg Val Asn Asp Ala Thr Ala Cys His
```

```
Ala Cys Lys Asn Val Tyr His Lys Gly Lys Cys Ile Glu Lys Cys Asp
Ala His Leu Tyr Leu Leu Gln Arg Arg Cys Val Thr Arg Glu Gln
                    70
Cys Leu Gln Leu Asn Pro Val Leu Ser Asn Lys Thr Val Pro Ile Lys
                                    90
               85
Ala Thr Ala Gly Leu Cys Ser Asp Lys Cys Pro Asp Gly Tyr Gln Ile
                               105
           100
                                                    110
Asn Pro Asp Asp His Arg Glu Cys Arg Lys Cys Val Gly Lys Cys Glu
                            120
       115
Ile Val Cys
   130
<210> 35
<211> 103
<212> PRT
<213> Caenorhabditis elegans
<400> 35
Phe Asp Gln Lys Ala Cys Glu Ser Leu Val Lys Lys Leu Lys Asp Lys
                                    10
Lys Asn Asp Leu Gln Asn Leu Ile Asp Val Val Leu Ser Lys Gly Thr
           20
                                25
Lys Tyr Thr Gly Cys Ile Thr Ile Pro Arg Thr Leu Asp Gly Arg Leu
                            40
Gln Val His Gly Arg Lys Gly Phe Pro His Val Val Tyr Gly Lys Leu
                        55
Trp Arg Phe Asn Glu Met Thr Lys Asn Glu Thr Arg His Val Asp His
                                        75
                    70
Cys Lys His Ala Phe Glu Met Lys Ser Asp Met Val Cys Val Asn Pro
                85
Tyr His Tyr Glu Ile Val Ile
           100
<210> 36
<211> 79
<212> PRT
<213> Caenorhabditis elegans
<400> 36
Asn Arg Tyr Ser Leu Gly Leu Glu Pro Asn Pro Ile Arg Glu Pro Val
1
                                    10
Ala Phe Lys Val Arg Lys Ala Ile Val Asp Gly Ile Arg Phe Ser Tyr
            2.0
                                25
Lys Lys Asp Gly Ser Val Trp Leu Gln Asn Arg Met Lys Tyr Pro Val
```

35 40 45 Phe Val Thr Ser Gly Tyr Leu Asp Glu Gln Ser Gly Gly Leu Lys Lys

Asp Lys Val His Lys Val Tyr Gly Cys Ala Ser Ile Lys Thr Phe

55

<210> 37

<211> 106

<212> PRT

<213> Caenorhabditis elegans

```
<400> 37
Lys Lys Thr Thr Thr Arg Arg Asn Ala Trp Gly Asn Met Ser Tyr Ala
                 5
                                    10
Glu Leu Ile Thr Thr Ala Ile Met Ala Ser Pro Glu Lys Arg Leu Thr
            20
                                25
Leu Ala Gln Val Tyr Glu Trp Met Val Gln Asn Val Pro Tyr Phe Arg
                            40
Asp Lys Gly Asp Ser Asn Ser Ser Ala Gly Trp Lys Asn Ser Ile Arg
    50
                        55
                                             60
His Asn Leu Ser Leu His Ser Arg Phe Met Arg Ile Gln Asn Glu Gly
                    70
                                        75
Ala Gly Lys Ser Ser Trp Trp Val Ile Asn Pro Asp Ala Lys Pro Gly
                85
                                    90
Met Asn Pro Arg Arg Thr Arg Glu Arg Ser
            100
<210> 38
```

<211> 60 <212> PRT <213> Caenorhabditis elegans

<210> 39 <211> 2784 <212> DNA <213> Caenorhabditis elegans

<400> 39

atgaagctaa tagcaacttc tcttctagtt cccgacgagc acacaccgat gatgtcacca 60 gtgaatacaa ctacaaagat tctacaacgg agtggtatta aaatggaaat cccgccatat 120 ttggatccag acagtcagga tgatgacccg gaagatggtg tcaactaccc ggatccagat 180 ttatttgaca caaaaaacac aaatatgacc gagtacgatt tggatgtgtt gaagcttgga 240 aaaccagcag tagatgaagc acggaaaaag atcgaagttc ccgacgctag tgcgccgcca 300 aacaaaattg tagaatattt gatgtattat agaacgttaa aagaaagtga actcatacaa 360 ctgaatgcgt atcggacaaa acgaaatcga ttatcgttga acttggtcaa aaacaatatt 420 gatcgagagt tcgaccaaaa agcttgcgag tccctggtga aaaaattgaa ggataagaag 480 aatgatetee agaacetgat tgatgtggtt ettteaaaag gtacaaaata taceggttge 540 attacaattc caaggacact tgatggccgg ttacaggtcc acggaagaaa aggtttccct 600 gtggaccact gcaagcacgc atttgaaatg aaaagtgaca tggtatgcgt gaatccctat 720 cactacgaaa ttgtcattgg aactatgatt gttgggcaga gggatcatga caatcgagat 780 atgccgccgc cacatcaacg ctaccacact ccaggtcggc aggatccagt tgacgatatg 840 agtagattta taccaccage ttecattegt eegecteega tgaacatgca cacaaggeet 900 cagoctatgc ctcaacaatt gccttcagtt ggcgcaacgt ttgcccatcc tctcccacat 960 caggegecae ataacceagg ggttteacat cegtaeteea ttgeteeaca gacceattae 1020 ccgttgaaca tgaacccaat tccgcaaatg ccgcaaatgc cacaaatgcc accacctctc 1080 catcagggat atggaatgaa tgggccgagt tgctcttcag aaaacaacaa tccattccac 1140

```
caaaatcacc attataatga tattagccat ccaaatcact attcctacga ctgtggtccg 1200
aacttgtacg ggtttccaac tccttatccg gattttcacc atcctttcaa tcagcaacca 1260
caccagoogc cacaactatc acaaaaccat acgtcccaac aaggcagtca tcaaccaggg 1320
caccaaggtc aggtaccgaa tgatccacca atttcaagac cagtgttaca accatcaaca 1380
gtcaccttgg acgtgttccg tcggtactgt agacagacat ttggaaatcg attttttgaa 1440
ggagaaagtg aacaatccgg cgcaataatt cggtctagta acaaattcat tgaagaattt 1500
gattcgccga tttgtggtgt gacagttgtt cgaccgcgga tgacagacgg tgaggttttg 1560
gagaacatca tgccggaaga tgcaccatat catgacattt gcaagttcat tttgaggctc 1620
acatcagaaa gtgtaacttt ctcaggagag gggccagaag ttagtgattt gaacgaaaaa 1680
tggggaacaa ttgtgtacta tgagaaaaat ttgcaaattg gcgagaaaaa atgttcgaga 1740
ggaaatttcc acgtggatgg cggattcatt tgctctgaga atcgttacag tctcggactt 1800
gagccaaatc caattagaga accagtggcg tttaaagttc gtaaagcaat agtggatgga 1860
attegetttt eetacaaaaa agaegggagt gtttggette aaaacegcat gaagtaceeg 1920
gtatttgtca cttctgggta tctcgacgag caatcaggag gcctaaagaa ggataaagtg 1980
cacaaagttt acggatgtgc gtctatcaaa acgtttggct tcaacgtttc caaacaaatc 2040
atcagagacg cgcttctttc caagcaaatg gcaacaatgt acttgcaagg aaaattgact 2100
ccgatgaatt atatctacga gaagaagact caggaagagc tgcgaaggga agcaacacgc 2160
accactgatt cattggccaa gtactgttgt gtccgtgtct cgttctgcaa aggatttgga 2220
gaaqcatacc cagaacgccc gtcaattcat gattgtccag tttggattga gttgaaaatc 2280
aacattgcct acgatttcat ggattcaatc tgccagtaca taaccaactg cttcgagccg 2340
ctaggaatgg aagattttgc aaaattggga atcaacgtca gtgatgacta aatgataact 2400
tttttcactc accctactag atactgattt agtcttattc caaatcatcc aacgatatca 2460
aactttttcc tttgaacttt gcatactatg ttatcacaag ttccaagcag tttcaataca 2520
aacataggat atgttaacaa cttttgataa gaatcaagtt accaactgtt cattgtgagc 2580
tttgagetgt atagaaggac aatgtateec ataceteaat etttaatagt cateagteac 2640
tggtcccgca ccaatttttt cgattcgcat atgtcatata ttgcaccgtg gcccttttta 2700
ttgtaacttt taatatattt tetteecaac ttgtgaatat gattgatgaa ceaccatttt 2760
                                                                  2784
gagtaataaa tgtatttttt gtgg
```

<210> 40 <211> 796

<212> PRT

<213> Caenorhabditis elegans

<400> 40

```
Met Lys Leu Ile Ala Thr Ser Leu Leu Val Pro Asp Glu His Thr Pro
                                    10
Met Met Ser Pro Val Asn Thr Thr Lys Ile Leu Gln Arg Ser Gly
                                25
Ile Lys Met Glu Ile Pro Pro Tyr Leu Asp Pro Asp Ser Gln Asp Asp
                            40
Asp Pro Glu Asp Gly Val Asn Tyr Pro Asp Pro Asp Leu Phe Asp Thr
                        55
                                             60
Lys Asn Thr Asn Met Thr Glu Tyr Asp Leu Asp Val Leu Lys Leu Gly
                    70
                                        75
Lys Pro Ala Val Asp Glu Ala Arg Lys Lys Ile Glu Val Pro Asp Ala
                85
                                    90
                                                         95
Ser Ala Pro Pro Asn Lys Ile Val Glu Tyr Leu Met Tyr Tyr Arg Thr
            100
                                105
                                                     110
Leu Lys Glu Ser Glu Leu Ile Gln Leu Asn Ala Tyr Arg Thr Lys Arg
                            120
                                                125
Asn Arg Leu Ser Leu Asn Leu Val Lys Asn Asn Ile Asp Arg Glu Phe
                        135
                                            140
Asp Gln Lys Ala Cys Glu Ser Leu Val Lys Lys Leu Lys Asp Lys Lys
                    150
                                        155
Asn Asp Leu Gln Asn Leu Ile Asp Val Val Leu Ser Lys Gly Thr Lys
                                    170
                165
Tyr Thr Gly Cys Ile Thr Ile Pro Arg Thr Leu Asp Gly Arg Leu Gln
```

			180					185					190		
Val	His	Glv		Tays	Glv	Phe	Pro		Val	Wa l	ጥህጉ	Glv		Leu	Trn
vai	1115	195	1119	Lys	OLY	1110	200	111.53	vai	var	1 <i>y</i> ±	205	шуы	шси	111
Arg	Phe 210	Asn	Glu	Met	Thr	Lys 215		Glu	Thr	Arg	His 220	Va1	Asp	His	Cys
Lys 225	His	Ala	Phe	Glu	Met 230	Lys	Ser	Asp	Met	Val 235	Сув	Val	Asn	Pro	Tyr 240
				245					250		-			Asp 255	
_		_	260					265					270	Pro	
_		275			_	_	280		_			285		Ala	
	290					295					300			Met	
305					310					315				Pro	320
				325					330					Ala 335	
			340					345					350	Pro	
		355					360					365		Asn	
	370					375					380			His	
385					390					395				Gly	400
				405					410					Pro 415	
			420					425					430	Thr	
		435					440					445		Asn	
	450					455					460			Leu	
465					470					475				Phe Lys	480
				485		_			490					495 Arg	
			500					505					510	Asp	
		515					520					525		Glu	
	530					535					540			Glu	
545					550					555				Glu	560
				565					570					575 Cys	
			580					585					590	Glu	
		595					600					605	_	Phe	
	610					615					620			Tyr	
625					630					635				Leu	640
val	r116	vaı	TIIT	per	σтλ	тАт	пеп	ASD	GIU	GTII	DGT.	атЛ	GTÀ	пец	пλр

```
645
                                   650
Lys Asp Lys Val His Lys Val Tyr Gly Cys Ala Ser Ile Lys Thr Phe
           660
                              665
Gly Phe Asn Val Ser Lys Gln Ile Ile Arg Asp Ala Leu Leu Ser Lys
                           680
Gln Met Ala Thr Met Tyr Leu Gln Gly Lys Leu Thr Pro Met Asn Tyr
                       695
                                           700
Ile Tyr Glu Lys Lys Thr Gln Glu Glu Leu Arg Arg Glu Ala Thr Arg
                                       715
                   710
Thr Thr Asp Ser Leu Ala Lys Tyr Cys Cys Val Arg Val Ser Phe Cys
                                   730
               725
Lys Gly Phe Gly Glu Ala Tyr Pro Glu Arg Pro Ser Ile His Asp Cys
           740
                               745
Pro Val Trp Ile Glu Leu Lys Ile Asn Ile Ala Tyr Asp Phe Met Asp
                           760
                                               765
Ser Ile Cys Gln Tyr Ile Thr Asn Cys Phe Glu Pro Leu Gly Met Glu
                       775
Asp Phe Ala Lys Leu Gly Ile Asn Val Ser Asp Asp
                   790
```

<210> 41 <211> 858 <212> PRT

<213> Caenorhabditis elegans

<400> 41 Met Gly Asp His His Asn Leu Thr Gly Leu Pro Gly Thr Ser Ile Pro 10 Pro Gln Phe Asn Tyr Ser Gln Pro Gly Thr Ser Thr Gly Gly Pro Leu 25 Tyr Gly Gly Lys Pro Ser His Gly Leu Glu Asp Ile Pro Asp Val Glu 40 Glu Tyr Glu Arg Asn Leu Leu Gly Ala Gly Ala Gly Phe Asn Leu Leu 55 60 Asn Val Gly Asn Met Ala Asn Val Pro Asp Glu His Thr Pro Met Met 70 75 Ser Pro Val Asn Thr Thr Lys Ile Leu Gln Arg Ser Gly Ile Lys 85 90 Met Glu Ile Pro Pro Tyr Leu Asp Pro Asp Ser Gln Asp Asp Pro 105 Glu Asp Gly Val Asn Tyr Pro Asp Pro Asp Leu Phe Asp Thr Lys Asn 120 Thr Asn Met Thr Glu Tyr Asp Leu Asp Val Leu Lys Leu Gly Lys Pro 135 140 Ala Val Asp Glu Ala Arg Lys Lys Ile Glu Val Pro Asp Ala Ser Ala 150 155 Pro Pro Asn Lys Ile Val Glu Tyr Leu Met Tyr Tyr Arg Thr Leu Lys 165 170 Glu Ser Glu Leu Ile Gln Leu Asn Ala Tyr Arg Thr Lys Arg Asn Arg 180 185 190 Leu Ser Leu Asn Leu Val Lys Asn Asn Ile Asp Arg Glu Phe Asp Gln 200 205 Lys Ala Cys Glu Ser Leu Val Lys Lys Leu Lys Asp Lys Lys Asn Asp 215 220 Leu Gln Asn Leu Ile Asp Val Val Leu Ser Lys Gly Thr Lys Tyr Thr 230 235 Gly Cys Ile Thr Ile Pro Arg Thr Leu Asp Gly Arg Leu Gln Val His

				245					250					255	
Gly	Arg	Lys	Gly 260	Phe	Pro	His	Val	Val 265	Tyr	Gly	Lys	Leu	Trp 270	Arg	Phe
Asn	Glu	Met 275	Thr	Lys	Asn	Glu	Thr 280	Arg	His	Val	Asp	His 285	Cys	Lys	His
Ala	Phe 290	Glu	Met	Lys	Ser	Asp 295	Met	Val	Cys	Val	Asn 300	Pro	Tyr	His	Tyr
Glu 305	Ile	Val	Ile	Gly	Thr 310	Met	Ile	Val	Gly	Gln 315	Arg	Asp	His	Asp	Asn 320
	_		Pro	325					330					335	
Asp	Pro	Val	Asp 340	Asp	Met	Ser	Arg	Phe 345	Ile	Pro	Pro	Ala	Ser 350	Ile	Arg
Pro	Pro	Pro 355	Met	Asn	Met	His	Thr 360	Arg	Pro	Gln	Pro	Met 365	Pro	Gln	Gln
	370		Val	_		375					380				
385			Pro		390					395					400
	_		Leu	405					410					415	
			Pro 420					425					430		
_		435	Glu				440					445			
	450		His			455					460				
465			Pro		470					475					480
			Gln	485					490					495	
-			Gln 500					505					510		
		515	Pro				520					525			
	530		Cys			535					540				
545	GIU	GIN	Ser	GTĀ	550	TTE	тте	Arg	ser	555	ASI	Lys	Pne	тте	560
Glu	Phe	Asp	Ser	Pro 565	Ile	Cys	Gly	Val	Thr 570	Val	Val	Arg	Pro	Arg 575	Met
Thr	Asp	Gly	Glu 580	Val	Leu	Glu	Asn	Ile 585	Met	Pro	Glu	Asp	Ala 590	Pro	Tyr
		595	Cys				600					605			
Phe	Ser 610	Gly	Glu	Gly	Pro	Glu 615	Val	Ser	Asp	Leu	Asn 620	Glu	Lys	Trp	Gly
Thr 625			Tyr	_	630					635	_		_	_	640
	_		Asn	645					650					655	
			Leu 660					665					670		
	_	675	Arg	_			680		_		-	685			
_	690	_	Ser			695					700				
Val	Thr	Ser	Gly	Tyr	Leu	Asp	Glu	Gln	Ser	Gly	Gly	Leu	Lys	Lys	Asp

willia th

```
705
                   710
                                       715
Lys Val His Lys Val Tyr Gly Cys Ala Ser Ile Lys Thr Phe Gly Phe
                                   730
               725
                                                        735
Asn Val Ser Lys Gln Ile Ile Arg Asp Ala Leu Leu Ser Lys Gln Met
                               745
           740
Ala Thr Met Tyr Leu Gln Gly Lys Leu Thr Pro Met Asn Tyr Ile Tyr
                            760
                                               765
Glu Lys Lys Thr Gln Glu Glu Leu Arg Arg Glu Ala Thr Arg Thr Thr
                       775
                                           780
Asp Ser Leu Ala Lys Tyr Cys Cys Val Arg Val Ser Phe Cys Lys Gly
                   790
                                       795
Phe Gly Glu Ala Tyr Pro Glu Arg Pro Ser Ile His Asp Cys Pro Val
               805
                                   810
Trp Ile Glu Leu Lys Ile Asn Ile Ala Tyr Asp Phe Met Asp Ser Ile
                               825
Cys Gln Tyr Ile Thr Asn Cys Phe Glu Pro Leu Gly Met Glu Asp Phe
       835
                           840
Ala Lys Leu Gly Ile Asn Val Ser Asp Asp
                        855
```

<210> 42 <211> 892 <212> PRT

<213> Caenorhabditis elegans

<400> 42 Met Gly Asp His His Asn Leu Thr Gly Leu Pro Gly Thr Ser Ile Pro Pro Gln Phe Asn Tyr Ser Gln Pro Gly Thr Ser Thr Gly Gly Pro Leu Tyr Gly Gly Lys Pro Ser His Gly Leu Glu Asp Ile Pro Asp Val Glu Glu Tyr Glu Arg Asn Leu Leu Gly Ala Gly Ala Gly Phe Asn Leu Leu Asn Val Gly Asn Met Ala Asn Glu Phe Lys Pro Ile Ile Thr Leu Asp Thr Lys Pro Pro Arg Asp Ala Asn Lys Ser Leu Ala Phe Asn Gly Gly Leu Lys Leu Ile Thr Pro Lys Thr Glu Val Pro Asp Glu His Thr Pro Met Met Ser Pro Val Asn Thr Thr Lys Ile Leu Gln Arg Ser Gly Ile Lys Met Glu Ile Pro Pro Tyr Leu Asp Pro Asp Ser Gln Asp Asp Asp Pro Glu Asp Gly Val Asn Tyr Pro Asp Pro Asp Leu Phe Asp Thr Lys Asn Thr Asn Met Thr Glu Tyr Asp Leu Asp Val Leu Lys Leu Gly Lys Pro Ala Val Asp Glu Ala Arg Lys Lys Ile Glu Val Pro Asp Ala Ser Ala Pro Pro Asn Lys Ile Val Glu Tyr Leu Met Tyr Tyr Arg Thr Leu Lys Glu Ser Glu Leu Ile Gln Leu Asn Ala Tyr Arg Thr Lys Arg Asn Arg Leu Ser Leu Asn Leu Val Lys Asn Asn Ile Asp Arg Glu Phe Asp Gln Lys Ala Cys Glu Ser Leu Val Lys Lys Leu Lys Asp Lys Lys

				245					250					255	
Asn	Asp	Leu	Gln 260	Asn	Leu	Ile	Asp	Val 265	Val	Leu	Ser	Lys	Gly 270	Thr	Lys
Tyr	Thr	Gly 275	Cys	Ile	Thr	Ile	Pro 280	Arg	Thr	Leu	Asp	Gly 285	Arg	Leu	Gln
Val	His 290	Gly	Arg	Lys	Gly	Phe 295	Pro	His	Val	Val	Tyr 300	Gly	Lys	Leu	Trp
Arg 305	Phe	Asn	Glu	Met	Thr 310	Lys	Asn	Glu	Thr	Arg 315	His	Val	Asp	His	Cys 320
				325	Met				330					335	
	_		340		Ile			345					350		
		355			Pro		360					365			
	370				Asp	375					380				
385	_				Met 390					395					400
				405	Val				410					415	
			420		Pro			425					430		
		435			Leu		440					445			
	450				Pro	455					460				
465					Glu 470					475					480
-				485	His				490					495	
			500		Pro Gln			505					510		
		515			Gln		520					525			
	530				Pro	535					540				
545					550 Cys					555					560
				565	Ser				570					575	
			580		Ser			585					590		
		595			Glu		600					605			
	610				Cys	615					620				
625	-1-				630	-1-				635					640
Val	Thr	Phe	Ser	Gly 645	Glu	Gly	Pro	Glu	Val 650	Ser	Asp	Leu	Asn	Glu 655	Lys
_	_		660		Tyr			665					670		
_	_	675		_	Asn		680				_	685			
	690				Leu	695					700				
Val	Ala	Phe	Lys	Val	Arg	Lys	Ala	Ile	Val	Asp	GГĀ	Ile	Arg	Phe	Ser

```
720
                    710
                                        715
705
Tyr Lys Lys Asp Gly Ser Val Trp Leu Gln Asn Arg Met Lys Tyr Pro
                                                        735
                                    730
                725
Val Phe Val Thr Ser Gly Tyr Leu Asp Glu Gln Ser Gly Gly Leu Lys
                                                    750
                                745
            740
Lys Asp Lys Val His Lys Val Tyr Gly Cys Ala Ser Ile Lys Thr Phe
                                                765
                            760
Gly Phe Asn Val Ser Lys Gln Ile Ile Arg Asp Ala Leu Leu Ser Lys
                        775
Gln Met Ala Thr Met Tyr Leu Gln Gly Lys Leu Thr Pro Met Asn Tyr
                                                             800
                                        795
                    790
Ile Tyr Glu Lys Lys Thr Gln Glu Glu Leu Arg Arg Glu Ala Thr Arg
                                    810
                805
Thr Thr Asp Ser Leu Ala Lys Tyr Cys Cys Val Arg Val Ser Phe Cys
                                825
            820
Lys Gly Phe Gly Glu Ala Tyr Pro Glu Arg Pro Ser Ile His Asp Cys
                                                 845
                            840
Pro Val Trp Ile Glu Leu Lys Ile Asn Ile Ala Tyr Asp Phe Met Asp
                                             860
                        855
Ser Ile Cys Gln Tyr Ile Thr Asn Cys Phe Glu Pro Leu Gly Met Glu
                                         875
                    870
Asp Phe Ala Lys Leu Gly Ile Asn Val Ser Asp Asp
                885
<210> 43
<211> 3499
<212> DNA
<213> Caenorhabditis elegans
<400> 43
tgatctttca agccgaagca atcaagacct caaagccaat caactctact cacttttctt 60
cagaacctta actttttgtg tcactttccc caaaaaccgt tcaagctgct gccttcactc 120
teatececte etettaetee ttetteteg teegetaeta etgtatette tggacateta 180
cetgtataca caccagtggc cagtcatctg ccattacaat ttcatcaatt gacacttctt 240
caacaacaac cgccgtcctc attcactccc gattcttcct catcctcaac atcgtcgtct 300
ttggctgaaa ttcccgaaga cgttatgatg gagatgctgg tagatcaggg aactgatgca 360
tegtcatecg cetecacgte caceteatet gtttegagat teggagegga caegtteatg 420
aatacaccgg atgatgtgat gatgaatgat gatatggaac cgattcctcg tgatcggtgc 480
aatacgtggc caatgcgtag gccgcaactc gaaccaccac tcaactcgag tcccattatt 540
catgaacaaa ttcctgaaga agatgctgac ctatacggga gcaatgagca atgtggacag 600
ctcggcggag catcttcaaa cgggtcgaca gcaatgcttc atactccaga tggaagcaat 660
teteateaga categtttet teggagttte agaatgteeg aategeeaga egatacegta 720
tcgggaaaaa agacaacgac cagacggaac gcttggggaa atatgtcata tgctgaactt 780
atcactacag ccattatggc tagtccagag aaacggttaa ctcttgcaca agtttacgaa 840
tggatggtcc agaatgttcc atacttcagg gataagggag attcgaacag ttcagctgga 900
 tggaagaact cgatccgtca caatctgtct cttcattctc gtttcatgcg aattcagaat 960
 gaaggagccg gaaagagctc gtggtgggtt attaatccag atgcaaagcc aggaatgaat 1020
 ccacggcgta cacgtgaacg atccaatact attgagacga ctacaaaggc tcaactcgaa 1080
 aaatctcgcc gcggagccaa gaagaggata aaggagagag cattgatggg ctcccttcac 1140
 tcgacactta atggaaattc gattgccgga tcgattcaaa cgatttctca cgatttgtat 1200
 gatgatgatc aatgcaagga gcatttgata acgttccatc atctttccgt ccccgaactc 1260
 aatcgaacct ctcgattcct ggatcgtcgt ctcgtgtttc tccagctatt ggaagtgata 1320
 tctatgatga tctagaattc ccatcatggg ttggcgaatc ggttccagca attccaagtg 1380
 atattgttga tagaactgat caaatgcgta tcgatgcaac tactcatagt tggtggagtt 1440
 cagattaagc aggagtcgaa gccgattaag acggaaccaa ttgctccacc accatcatac 1500
```

cacgagttga acagtgtccg tggatcgtgt gctcagaatc cacttcttcg aaatccaatt 1560 gtgccaagca ctaacttcaa gccaatgcca ctaccgggtg cctatggaaa ctatcaaaat 1620

```
ggtggaataa ctccaatcaa ttggctatca acatccaact catctccact gcctggaatt 1680
caatcgtgtg gaattgtagc tgcacagcat actgtcgctt cttcatcggc tcttccaatt 1740
gatttggaaa atctgacact tcccgatcag ccactgatgg atactatgga tgttgatgca 1800
ttgatcagac atgagctgag tcaagctgga gggcagcata ttcattttga tttgtaaatt 1860
ctcttcattt tgtttcccct ggtgttgttc gaaagagaga tagcaaagca gcgaggagtg 1920
aggtaagcag caataaaaat tttggatttt tttttggttt ttccagaaat aatcgatttt 1980
ctggaaaatt tcaaaaaaaa atcggaattt ttagttaatt atttgatgag aaaaaaaaat 2040
tagaaaacat aaggaaaaat gaaaagcgtt tttttttttc gaaaatttta gaattctcct 2100
acatttccaa taagggcctt agaactgcaa acaaacaaaa attggaattt tcgaatcaaa 2160
aagttcccga ataaaagtag ttcgaatatt aaaaagcatt taatttcctc tttaaaaaaa 2220
ttgaataata gccgaaattt gcagattttt tttctgaaaa tcgaaaaacc aaaatttttt 2280
gattttttaa atttttttt tactttccag atagtaaaat cattagcact gaaaattatt 2340
tgaaaaaaaa cttcaaatac aaattttgtt ttcgaaaaaa aaaatttaaa tatatatttt 2400
cagaaatctt ccgtcttcat cttttcaaat ccctacctac acacactcaa cgatcatcac 2460
agccagacca tcaatattct tccaaatttt gacgtcgtta attttttttc agttttttca 2520
aaaactctat tttctatttt ctgtcgtttg ttcccctttc tctcgtctaa ttccaacaca 2580
ttcatcccag tgacgtcgtg taataataat ataaaatacc tcttctctct ttcttcccct 2640
aatgcgaaat atcgaaaaac cgttgattat tacctctttt ttcttgtttt ttttttctct 2700
ctctctctcc cgtcatccag gttcttcact ctttaaatgc tacctctatc ccatcttttt 2760
cgctgtaaat ttgtttcgca atcaaaactg ctaaaacaca ttccccaatc tgtctttttt 2820
aattgaattt ticaaaaaat tigattictt gattictctt gtaatticttt aattiticctc 2880
ttttttttcc ccctggtagc aaatgtctag cgattctctt tctttttttg tttaactttc 2940
acatctggcc gattcgaatc ctccgtatac acacacat agtaatctac ctccaaaatt 3000
ttactgaaag atgtgatccc ctctctgtct ccctctacaa aacattattt gtctgtttgt 3060
gtatattgcc accacgtcga ttttaaatta aaaccatcgt tttttcttct tttctacttt 3120
tttctcgaaa aatttaacaa cacacaaaaa aatccttcaa aaaatctcag ttttaaatgg 3180
tgtggcaata tatcggatcc ccctctacac cagaacagtc ttgcaatttc agagaatgat 3240
tttcagattt ttcatatcac aggccccctt tttttgcttg ttttttctc tacctctctt 3300
tottttcatt ctatttctct ctcttgtttt ctctctgtta tcctgtacat tttccttcca 3360
attettete getatttete attttegagt teatattete taegteteae tttetetege 3420
gccacgcccc ctttttcgtc tccctccgcc cccaaatata tttgcgactg tatgatgatg 3480
                                                                  3499
atgatgattt aataaaaat
<210> 44
<211> 2704
<212> DNA
<213> Caenorhabditis elegans
<400> 44
ttacacgtgg ccaatgcaac aatacatcta tcaggaatcg tcagcaacca ttccccatca 60
ccatttaaat caacacaaca atccgtatca tccaatgcat cctcatcatc aattacctca 120
tatgcaacaa cttcctcaac ctctattgaa tcttaacatg acgacgttaa catcttctgg 180
cagttccgtg gccagttcca ttggaggcgg agctcaatgc tctccgtgcg cgtcgggctc 240
ctcgaccgct gcaacaaatt cctctcaaca gcagcagacc gttggtcaaa tgcttgctgc 300
ateggtgeet tgttetteat etggeatgae aettggaatg teaettaate tgteaeaagg 360
cggtggtcca atgccggcaa aaaagaagcg ttgtcgtaag aagccaaccg atcaattggc 420
acagaagaaa ccgaatccat ggggtgagga atcctattcg gatatcattg ccaaagcatt 480
ggaatcggcg ccagacggaa ggcttaaact caatgagatt tatcaatggt tctctgataa 540
tattccctac titggagaac gatctagtcc cgaggaggcc gccggatgga agaactcgat 600
ccgtcacaat ctgtctcttc attctcgttt catgcgaatt cagaatgaag gagccggaaa 660
gagctcgtgg tgggttatta atccagatgc aaagccagga atgaatccac ggcgtacacg 720
tgaacgatcc aatactattg agacgactac aaaggctcaa ctcgaaaaat ctcgccgcgg 780
agccaagaag aggataaagg agagagcatt gatgggctcc cttcactcga cacttaatgg 840
aaattegatt geeggatega ticaaaegat tieteaegat tigtatgatg atgatteaat 900
gcaaggagca tttgataacg ttccatcatc tttccgtccc cgaactcaat cgaacctctc 960
gatteetgga tegtegtete gtgtttetee agetattgga agtgatatet atgatgatet 1020
```

agaattccca tcatgggttg gcgaatcggt tccagcaatt ccaagtgata ttgttgatag 1080 aactgatcaa atgcgtatcg atgcaactac tcatattggt ggagttcaga ttaagcagga 1140

```
gtcgaagccg attaagacgg aaccaattgc tecaccacca tcataccacg agttgaacag 1200
tgtccgtgga tcgtgtgctc agaatccact tcttcgaaat ccaattgtgc caagcactaa 1260
cttcaagcca atgccactac cgggtgccta tggaaactat caaaatggtg gaataactcc 1320
aatcaattgg ctatcaacat ccaactcatc tccactgcct ggaattcaat cgtgtggaat 1380
tgtagctgca cagcatactg tcgcttcttc atcggctctt ccaattgatt tggaaaatct 1440
gacacttccc gatcagccac tgatggatac tatggatgtt gatgcattga tcagacatga 1500
gctgagtcaa gctggagggc agcatattca ttttgatttg taaattctct tcattttgtt 1560
teceetggtg ttgttegaaa gagagatage aaageagega ggagtgagaa atetteegte 1620
ttcatctttt caaatcccta cctacacaca ctcaacgatc atcacagcca gaccatcaat 1680
attetteeaa attitgaegt egitaattit titteagtit titteaaaaae tetattitet 1740
attitictgtc gtttgttccc ctttctctcg tctaattcca acacattcat cccagtgacg 1800
tegtgtaata ataatataaa ataeetette tetetttett eecetaatge gaaatatega 1860
aaaaccgttg attattacct cttttttctt gtttttttt tctctctctc tctcccgtca 1920
tccaggttct tcactcttta aatgctacct ctatcccatc tttttcgctg taaatttgtt 1980
tegeaateaa aactgetaaa acaeatteee caatetgtet titttaattg aattitteaa 2040
aaaatttgat ttcttgattt ctcttgtaat tctttaattt tcctctttt tttccccctg 2100
gtagcaaatg tctagcgatt ctctttcttt ttttgtttaa ctttcacatc tggccgattc 2160
gaatcctccg tatacacaca cacatagtaa tctacctcca aaattttact gaaagatgtg 2220
atcccctctc tgtctccctc tacaaaacat tatttgtctg tttgtgtata ttgccaccac 2280
gtcgatttta aattaaaacc atcgtttttt cttcttttct actttttct cgaaaaattt 2340
aacaacacac aaaaaaatcc ttcaaaaaat ctcagtttta aatggtgtgg caatatatcg 2400
gatccccctc tacaccagaa cagtcttgca atttcagaga atgattttca gatttttcat 2460
atcacaggcc cccttttttt gcttgttttt ttctctacct ctctttcttt tcattctatt 2520
tetetetett gttttetete tgttateetg taeattttee tteeaattet ttetggetat 2580
ttctgatttt cgagttcata ttctctacgt ctcactttct ctcgcgccac gccccctttt 2640
tegteteeet eegeeeeeaa atatatttge gaetgtatga tgatgatgat gatttaataa 2700
<210> 45
<211> 510
<212> PRT
<213> Caenorhabditis elegans
<400> 45
Met Met Glu Met Leu Val Asp Gln Gly Thr Asp Ala Ser Ser Ser Ala
                                    10
```

Ser Thr Ser Thr Ser Ser Val Ser Arg Phe Gly Ala Asp Thr Phe Met 20 25 Asn Thr Pro Asp Asp Val Met Met Asn Asp Asp Met Glu Pro Ile Pro 40 45 Arg Asp Arg Cys Asn Thr Trp Pro Met Arg Arg Pro Gln Leu Glu Pro 55 60 Pro Leu Asn Ser Ser Pro Ile Ile His Glu Gln Ile Pro Glu Glu Asp 70 75 Ala Asp Leu Tyr Gly Ser Asn Glu Gln Cys Gly Gln Leu Gly Gly Ala 85 95 90 Ser Ser Asn Gly Ser Thr Ala Met Leu His Thr Pro Asp Gly Ser Asn 100 105 110 Ser His Gln Thr Ser Phe Pro Ser Asp Phe Arg Met Ser Glu Ser Pro 120 125 Asp Asp Thr Val Ser Gly Lys Lys Thr Thr Thr Arg Arg Asn Ala Trp 130 135 140 Gly Asn Met Ser Tyr Ala Glu Leu Ile Thr Thr Ala Ile Met Ala Ser 150 155 Pro Glu Lys Arg Leu Thr Leu Ala Gln Val Tyr Glu Trp Met Val Gln 165 170 1.75 Asn Val Pro Tyr Phe Arg Asp Lys Gly Asp Ser Asn Ser Ser Ala Gly 180 185

```
Trp Lys Asn Ser Ile Arg His Asn Leu Ser Leu His Ser Arg Phe Met
                           200
       195
Arg Ile Gln Asn Glu Gly Ala Gly Lys Ser Ser Trp Trp Val Ile Asn
                      215
                                          220
Pro Asp Ala Lys Pro Gly Met Asn Pro Arg Arg Thr Arg Glu Arg Ser
                  230
                                      235
Asn Thr Ile Glu Thr Thr Lys Ala Gln Leu Glu Lys Ser Arg Arg
              245
                       250
Gly Ala Lys Lys Arg Ile Lys Glu Arg Ala Leu Met Gly Ser Leu His
                              265
          260
Ser Thr Leu Asn Gly Asn Ser Ile Ala Gly Ser Ile Gln Thr Ile Ser
                           280
                                              285
His Asp Leu Tyr Asp Asp Ser Met Gln Gly Ala Phe Asp Asn Val
                      295
                                         300
Pro Ser Ser Phe Arg Pro Arg Thr Gln Ser Asn Leu Ser Ile Pro Gly
                  310
                                      315
Ser Ser Ser Arg Val Ser Pro Ala Ile Gly Ser Asp Ile Tyr Asp Asp
               325
                                  330
Leu Glu Phe Pro Ser Trp Val Gly Glu Ser Val Pro Ala Ile Pro Ser
                              345
Asp Ile Val Asp Arg Thr Asp Gln Met Arg Ile Asp Ala Thr Thr His
                           360
                                             365
Ile Gly Gly Val Gln Ile Lys Gln Glu Ser Lys Pro Ile Lys Thr Glu
                      375
                                          380
Pro Ile Ala Pro Pro Pro Ser Tyr His Glu Leu Asn Ser Val Arg Gly
                  390
                                      395
Ser Cys Ala Gln Asn Pro Leu Leu Arg Asn Pro Ile Val Pro Ser Thr
               405
                                 410
Asn Phe Lys Pro Met Pro Leu Pro Gly Ala Tyr Gly Asn Tyr Gln Asn
           420
                    425
                                                 430
Gly Gly Ile Thr Pro Ile Asn Trp Leu Ser Thr Ser Asn Ser Ser Pro
                          440
                                   445
Leu Pro Gly Ile Gln Ser Cys Gly Ile Val Ala Ala Gln His Thr Val
                       455
                                          460
Ala Ser Ser Ser Ala Leu Pro Ile Asp Leu Glu Asn Leu Thr Leu Pro
                  470
                                     475
Asp Gln Pro Leu Met Asp Thr Met Asp Val Asp Ala Leu Ile Arg His
              485
                                  490
Glu Leu Ser Gln Ala Gly Gly Gln His Ile His Phe Asp Leu
           500
                               505
```

```
<210> 46
```

<400> 46

 Met
 Gln
 Gln
 Tyr
 Ile
 Tyr
 Gln
 Glu
 Ser
 Ala
 Thr
 Ile
 Pro
 His
 Pro
 His
 Pro
 Met
 His
 Pro
 Met
 His
 Pro
 His
 Pro
 His
 Pro
 Leu
 His
 Pro
 His
 His
 Pro
 His
 His
 Pro
 His
 Pro
 Met
 His
 Pro
 Leu
 Pro
 Leu
 Leu
 Asn
 Leu
 Asn
 A

<211> 509

<212> PRT

<213> Caenorhabditis elegans

```
Thr Asn Ser Ser Gln Gln Gln Gln Thr Val Gly Gln Met Leu Ala Ala
                                  90
Ser Val Pro Cys Ser Ser Ser Gly Met Thr Leu Gly Met Ser Leu Asn
                              105
           100
Leu Ser Gln Gly Gly Pro Met Pro Ala Lys Lys Lys Arg Cys Arg
                          120
Lys Lys Pro Thr Asp Gln Leu Ala Gln Lys Lys Pro Asn Pro Trp Gly
                                         140
                      135
Glu Glu Ser Tyr Ser Asp Ile Ile Ala Lys Ala Leu Glu Ser Ala Pro
                                     155
       150
Asp Gly Arg Leu Lys Leu Asn Glu Ile Tyr Gln Trp Phe Ser Asp Asn
                               170
            165
Ile Pro Tyr Phe Gly Glu Arg Ser Ser Pro Glu Glu Ala Ala Gly Trp
                                                 190
                              185
Lys Asn Ser Ile Arg His Asn Leu Ser Leu His Ser Arg Phe Met Arg
                          200
                                             205
       195
Ile Gln Asn Glu Gly Ala Gly Lys Ser Ser Trp Trp Val Ile Asn Pro
                                          220
                      215
Asp Ala Lys Pro Gly Met Asn Pro Arg Arg Thr Arg Glu Arg Ser Asn
                        235
                 230
Thr Ile Glu Thr Thr Lys Ala Gln Leu Glu Lys Ser Arg Arg Gly
                                 250
               245
Ala Lys Lys Arg Ile Lys Glu Arg Ala Leu Met Gly Ser Leu His Ser
                             265
Thr Leu Asn Gly Asn Ser Ile Ala Gly Ser Ile Gln Thr Ile Ser His
                                              285
                          280
Asp Leu Tyr Asp Asp Asp Ser Met Gln Gly Ala Phe Asp Asn Val Pro
                                          300
                       295
Ser Ser Phe Arg Pro Arg Thr Gln Ser Asn Leu Ser Ile Pro Gly Ser
                                      315
                   310
Ser Ser Arg Val Ser Pro Ala Ile Gly Ser Asp Ile Tyr Asp Asp Leu
                                 330
               325
Glu Phe Pro Ser Trp Val Gly Glu Ser Val Pro Ala Ile Pro Ser Asp
                                                  350
                               345
            340
Ile Val Asp Arg Thr Asp Gln Met Arg Ile Asp Ala Thr Thr His Ile
                                              365
                           360
        355
Gly Gly Val Gln Ile Lys Gln Glu Ser Lys Pro Ile Lys Thr Glu Pro
                                    380
                       375
    370
Ile Ala Pro Pro Pro Ser Tyr His Glu Leu Asn Ser Val Arg Gly Ser
                                       395
                   390
Cys Ala Gln Asn Pro Leu Leu Arg Asn Pro Ile Val Pro Ser Thr Asn
                                   410
                405
Phe Lys Pro Met Pro Leu Pro Gly Ala Tyr Gly Asn Tyr Gln Asn Gly
                               425
            420
Gly Ile Thr Pro Ile Asn Trp Leu Ser Thr Ser Asn Ser Ser Pro Leu
                           440
Pro Gly Ile Gln Ser Cys Gly Ile Val Ala Ala Gln His Thr Val Ala
                        455
Ser Ser Ser Ala Leu Pro Ile Asp Leu Glu Asn Leu Thr Leu Pro Asp
                                       475
                    470
Gln Pro Leu Met Asp Thr Met Asp Val Asp Ala Leu Ile Arg His Glu
                                   490
                485
Leu Ser Gln Ala Gly Gly Gln His Ile His Phe Asp Leu
                               505
```

<210> 47 <211> 3504

cggaagccat ggagctcgag atctgattgc tggacacgga cggaactccg acgtatctcg 60 cagatgcatg ttaacatttt acatccacaa ctgcaaacga tggtcgagca gtggcaaatg 120 cgagaacgcc catcgctgga gaccgagaat ggcaaaggat cgctgctcct ggaaaatgaa 180 ggtgtcgcag atatcatcac tatgtgtcca ttcggagaag ttattagtgt agtatttccg 240 tggtttcttg caaatgtgcg aacatcgcta gaaatcaagc tatcagattt caaacatcaa 300 cttttcgaat tgattgctcc gatgaagtgg ggaacatatt ccgtaaagcc acaggattat 360 gtgttcagac agttgaataa tttcggcgaa attgaagtta tatttaacga cgatcaaccc 420 ctgtcgaaat tagagctcca cggcactttc ccaatgcttt ttctctacca acctgatgga 480 ataaacaggg ataaagaatt aatgagtgat ataagtcatt gtctaggata ctcactggat 540 aaactggaag agagcctcga tgaggaactc cgtcaatttc gtgcttctct ctgggctcgt 600 acgaagaaaa cgtgcttgac acgtggactt gagggtacca gtcactacgc gttccccgaa 660 gaacagtact tgtgtgttgg tgaatcgtgc ccgaaagatt tggaatcaaa agtcaaggct 720 gccaagctga gttatcagat gttttggaga aaacgtaaag cggaaatcaa tggagtttgc 780 gagaaaatga tgaagattca aattgaattc aatccgaacg aaactccgaa atctctgctt 840 cacacgtttc tctacgaaat gcgaaaattg gatgtatacg ataccgatga tcctgcagat 900 gaaggatggt ttcttcaatt ggctggacgt accacgtttg ttacaaatcc agatgtcaaa 960 cttacgtctt atgatggtgt ccgttcggaa ctggaaagct atcgatgccc tggattcgtt 1020 gttcgccgac aatcactagt cctcaaagac tattgtcgcc caaaaccact ctacgaacca 1080 cattatgtga gagcacacga acgaaaactt gctctagacg tgctcagcgt gtctatagat 1140 agcacaccaa aacagagcaa gaacagtgac atggttatga ctgattttcg tccgacagct 1200 tcactcaaac aagtttcact ttgggacctt gacgcgaatc ttatgatacg gcctgtgaat 1260 atttctggat tcgatttccc ggccgacgtg gatatgtacg ttcgaatcga attcagtgta 1320 tatgtgggga cactgacgct ggcatcaaaa tctacaacaa aagtgaatgc tcaatttgca 1380 aaatggaata aggaaatgta cacttttgat ctatacatga aggatatgcc accatctgca 1440 gtactcagca ttcgtgtttt gtacggaaaa gtgaaattaa aaagtgaaga attcgaagtt 1500 ggttgggtaa atatgtccct aaccgattgg agagatgaac tacgacaagg acaattttta 1560 ttccatctgt gggctcctga accgactgcc aatcgtagta ggatcggaga aaatggagca 1620 aggataggca ccaacgcagc ggttacaatt gaaatctcaa gttatggtgg tagagttcga 1680 atgccgagtc aaggacaata cacatatctc gtcaagcacc gaagtacttg gacggaaact 1740 ttgaatatta tgggtgatga ctatgagtcg tgtatcagag atccaggata taagaagctt 1800 cagatgettg teaagaagea tgaatetgga attgtattag aggaagatga acaaegteat 1860 gtctggatgt ggaggagata cattcaaaag caggagcctg atttgctcat tgtgctctcc 1920 gaactcgcat ttgtgtggac tgatcgtgag aacttttccg agctctatgt gatgcttgaa 1980 aaatggaaac cgccgagtgt ggcagccgcg ttgactttgc ttggaaaacg ttgcacggat 2040 cgtgtgattc gaaagtttgc agtggagaag ttgaatgagc agctgagccc ggtcacattc 2100 catcttttca tattgcctct catacaggcg ttgaagtacg aaccgcgtgc tcaatcggaa 2160 gttggaatga tgctcttgac tagagctctc tgcgattatc gaattggaca tcgacttttc 2220 tggctgctcc gtgcagagat tgctcgtttg agagattgtg atctgaaaag tgaagaatat 2280 cgccgtatct cacttctgat ggaagcttac ctccgtggaa atgaagagca catcaagatc 2340 atcacccgac aagttgacat ggttgatgag ctcacacgaa tcagcactct tgtcaaagga 2400 atgccaaaag atgttgctac gatgaaactg cgtgacgagc ttcgatcgat tagtcataaa 2460 atggaaaata tggattctcc actggatcct gtgtacaaac tgggtgaaat gataatcgac 2520 aaagccatcg tcctaggaag tgcaaaacgt ccgttaatgc ttcactggaa gaacaaaaat 2580 ccaaagagtg acctgcacct tccgttctgt gcaatgatct tcaagaatgg agacgatctt 2640 cgccaggaca tgcttgttct tcaagttctc gaagttatgg ataacatctg gaaggctgca 2700 aacattgatt gctgtttgaa cccgtacgca gttcttccaa tgggagaaat gattggaatt 2760 attgaagttg tgcctaattg taaaacaata ttcgagattc aagttggaac aggattcatg 2820 aatacagcag ttcggagtat tgatccttcg tttatgaata agtggattcg gaaacaatgc 2880 ggaattgaag atgaaaagaa gaaaagcaaa aaggactcta cgaaaaatcc catcgaaaag 2940 aagattgata atactcaagc catgaagaaa tattttgaaa gtgtcgatcg attcctatac 3000 tcgtgtgttg gatattcagt tgccacgtac ataatgggaa tcaaggatcg tcacagtgat 3060 aatctgatgc tcactgaaga tggaaaatat gtccacattg atttcggtca cattttggga 3120 cacggaaaga ccaaacttgg gatccagcga gatcgtcaac cgtttattct aaccgaacac 3180 tttatgacag tgattcgatc gggtaaatct gtggatggaa attcgcatga gctacaaaaa 3240

```
ttcaaaacgt tatgcgtcga agcctacgaa gtaatgtgga ataatcgaga tttgttcgtt 3300
tccttgttca ccttgatgct cggaatggag ttgcctgagc tgtcgacgaa agcggatttg 3360
gatcatttga agaaaaccct cttctgcaat ggagaaagca aagaagaagc gagaaagttt 3420
ttcgctggaa tctacgaaga agccttcaat ggatcatggt ctaccaaaac gaattggctc 3480
ttccacgcag tcaaacacta ctga
<210> 48
<211> 1167
<212> PRT
<213> Caenorhabditis elegans
<400> 48
Arg Lys Pro Trp Ser Ser Arg Ser Asp Cys Trp Thr Arg Thr Glu Leu
                                    10
                 5
Arg Arg Ile Ser Gln Met His Val Asn Ile Leu His Pro Gln Leu Gln
                                                     30
                                25
            20
Thr Met Val Glu Gln Trp Gln Met Arg Glu Arg Pro Ser Leu Glu Thr
                            40
Glu Asn Gly Lys Gly Ser Leu Leu Leu Glu Asn Glu Gly Val Ala Asp
                        55
Ile Ile Thr Met Cys Pro Phe Gly Glu Val Ile Ser Val Val Phe Pro
                                         75
                    70
Trp Phe Leu Ala Asn Val Arg Thr Ser Leu Glu Ile Lys Leu Ser Asp
                                     90
                85
Phe Lys His Gln Leu Phe Glu Leu Ile Ala Pro Met Lys Trp Gly Thr
                                                     110
                                105
            100
Tyr Ser Val Lys Pro Gln Asp Tyr Val Phe Arg Gln Leu Asn Asn Phe
                            120
                                                 125
        115
Gly Glu Ile Glu Val Ile Phe Asn Asp Asp Gln Pro Leu Ser Lys Leu
                                             140
Glu Leu His Gly Thr Phe Pro Met Leu Phe Leu Tyr Gln Pro Asp Gly
                                        155
                    150
Ile Asn Arg Asp Lys Glu Leu Met Ser Asp Ile Ser His Cys Leu Gly
                                                         175
                                     170
                165
Tyr Ser Leu Asp Lys Leu Glu Glu Ser Leu Asp Glu Glu Leu Arg Gln
                                                     190
                                 185
            180
Phe Arg Ala Ser Leu Trp Ala Arg Thr Lys Lys Thr Cys Leu Thr Arg
                                                 205
                             200
        195
Gly Leu Glu Gly Thr Ser His Tyr Ala Phe Pro Glu Glu Gln Tyr Leu
                                             220
                         215
    210
Cys Val Gly Glu Ser Cys Pro Lys Asp Leu Glu Ser Lys Val Lys Ala
                                         235
                     230
225
Ala Lys Leu Ser Tyr Gln Met Phe Trp Arg Lys Arg Lys Ala Glu Ile
                                     250
                 245
Asn Gly Val Cys Glu Lys Met Met Lys Ile Gln Ile Glu Phe Asn Pro
                                 265
Asn Glu Thr Pro Lys Ser Leu Leu His Thr Phe Leu Tyr Glu Met Arg
                                                 285
                             280
Lys Leu Asp Val Tyr Asp Thr Asp Asp Pro Ala Asp Glu Gly Trp Phe
                                             300
                         295
Leu Gln Leu Ala Gly Arg Thr Thr Phe Val Thr Asn Pro Asp Val Lys
                                         315
                     310
Leu Thr Ser Tyr Asp Gly Val Arg Ser Glu Leu Glu Ser Tyr Arg Cys
                                     330
                 325
Pro Gly Phe Val Val Arg Arg Gln Ser Leu Val Leu Lys Asp Tyr Cys
```

365

345

Arg Pro Lys Pro Leu Tyr Glu Pro His Tyr Val Arg Ala His Glu Arg

360

340

355

```
Lys Leu Ala Leu Asp Val Leu Ser Val Ser Ile Asp Ser Thr Pro Lys
                                           380
                       375
Gln Ser Lys Asn Ser Asp Met Val Met Thr Asp Phe Arg Pro Thr Ala
                                      395
                390
Ser Leu Lys Gln Val Ser Leu Trp Asp Leu Asp Ala Asn Leu Met Ile
                                   410
               405
Arg Pro Val Asn Ile Ser Gly Phe Asp Phe Pro Ala Asp Val Asp Met
                               425
           420
Tyr Val Arg Ile Glu Phe Ser Val Tyr Val Gly Thr Leu Thr Leu Ala
                                               445
                           440
       435
Ser Lys Ser Thr Thr Lys Val Asn Ala Gln Phe Ala Lys Trp Asn Lys
                      455
                                          460
Glu Met Tyr Thr Phe Asp Leu Tyr Met Lys Asp Met Pro Pro Ser Ala
                                       475
                  470
Val Leu Ser Ile Arg Val Leu Tyr Gly Lys Val Lys Leu Lys Ser Glu
                                   490
               485
Glu Phe Glu Val Gly Trp Val Asn Met Ser Leu Thr Asp Trp Arg Asp
                                                   510
                            505
Glu Leu Arg Gln Gly Gln Phe Leu Phe His Leu Trp Ala Pro Glu Pro
                    520
Thr Ala Asn Arg Ser Arg Ile Gly Glu Asn Gly Ala Arg Ile Gly Thr
                                           540
                        535
Asn Ala Ala Val Thr Ile Glu Ile Ser Ser Tyr Gly Gly Arg Val Arg
                                       555
                    550
Met Pro Ser Gln Gly Gln Tyr Thr Tyr Leu Val Lys His Arg Ser Thr
                                  570
                565
Trp Thr Glu Thr Leu Asn Ile Met Gly Asp Asp Tyr Glu Ser Cys Ile
                               585
Arg Asp Pro Gly Tyr Lys Lys Leu Gln Met Leu Val Lys Lys His Glu
                                               605
                           600
Ser Gly Ile Val Leu Glu Glu Asp Glu Gln Arg His Val Trp Met Trp
                                          620
                        615
Arg Arg Tyr Ile Gln Lys Gln Glu Pro Asp Leu Leu Ile Val Leu Ser
                                      635
                    630
Glu Leu Ala Phe Val Trp Thr Asp Arg Glu Asn Phe Ser Glu Leu Tyr
                                    650
                645
Val Met Leu Glu Lys Trp Lys Pro Pro Ser Val Ala Ala Ala Leu Thr
                                665
            660
Leu Leu Gly Lys Arg Cys Thr Asp Arg Val Ile Arg Lys Phe Ala Val
                                                685
                            680
        675
Glu Lys Leu Asn Glu Gln Leu Ser Pro Val Thr Phe His Leu Phe Ile
                                            700
                        695
Leu Pro Leu Ile Gln Ala Leu Lys Tyr Glu Pro Arg Ala Gln Ser Glu
                                        715
                    710
Val Gly Met Met Leu Leu Thr Arg Ala Leu Cys Asp Tyr Arg Ile Gly
                                    730
                725
 His Arg Leu Phe Trp Leu Leu Arg Ala Glu Ile Ala Arg Leu Arg Asp
                                745
 Cys Asp Leu Lys Ser Glu Glu Tyr Arg Arg Ile Ser Leu Leu Met Glu
                            760
 Ala Tyr Leu Arg Gly Asn Glu Glu His Ile Lys Ile Ile Thr Arg Gln
                                            780
                        775
 Val Asp Met Val Asp Glu Leu Thr Arg Ile Ser Thr Leu Val Lys Gly
                                        795
                     790
 Met Pro Lys Asp Val Ala Thr Met Lys Leu Arg Asp Glu Leu Arg Ser
                                    810
 Ile Ser His Lys Met Glu Asn Met Asp Ser Pro Leu Asp Pro Val Tyr
                                825
```

```
Lys Leu Gly Glu Met Ile Ile Asp Lys Ala Ile Val Leu Gly Ser Ala
                     840
Lys Arg Pro Leu Met Leu His Trp Lys Asn Lys Asn Pro Lys Ser Asp
                                  860
            855
Leu His Leu Pro Phe Cys Ala Met Ile Phe Lys Asn Gly Asp Asp Leu
                               875
    870
Arg Gln Asp Met Leu Val Leu Gln Val Leu Glu Val Met Asp Asn Ile
                    890
          885
Trp Lys Ala Ala Asn Ile Asp Cys Cys Leu Asn Pro Tyr Ala Val Leu
        900
                       905
Pro Met Gly Glu Met Ile Gly Ile Ile Glu Val Val Pro Asn Cys Lys
           920
                                     925
Thr Ile Phe Glu Ile Gln Val Gly Thr Gly Phe Met Asn Thr Ala Val
       935
                                940
Arg Ser Ile Asp Pro Ser Phe Met Asn Lys Trp Ile Arg Lys Gln Cys
       950 955
Gly Ile Glu Asp Glu Lys Lys Lys Ser Lys Lys Asp Ser Thr Lys Asn
                                   975
            965 970
Pro Ile Glu Lys Lys Ile Asp Asn Thr Gln Ala Met Lys Lys Tyr Phe
                                      990
             985
Glu Ser Val Asp Arg Phe Leu Tyr Ser Cys Val Gly Tyr Ser Val Ala
                           1005
                     1000
Thr Tyr Ile Met Gly Ile Lys Asp Arg His Ser Asp Asn Leu Met Leu
                          1020
   1010
                  1015
Thr Glu Asp Gly Lys Tyr Val His Ile Asp Phe Gly His Ile Leu Gly
                     1035 1040
               1030
His Gly Lys Thr Lys Leu Gly Ile Gln Arg Asp Arg Gln Pro Phe Ile
                            1050 1055
            1045
Leu Thr Glu His Phe Met Thr Val Ile Arg Ser Gly Lys Ser Val Asp
         1060 1065 1070
Gly Asn Ser His Glu Leu Gln Lys Phe Lys Thr Leu Cys Val Glu Ala
      1075 1080 1085
Tyr Glu Val Met Trp Asn Asn Arg Asp Leu Phe Val Ser Leu Phe Thr
   1090 1095 1100
Leu Met Leu Gly Met Glu Leu Pro Glu Leu Ser Thr Lys Ala Asp Leu
      1110 1115
Asp His Leu Lys Lys Thr Leu Phe Cys Asn Gly Glu Ser Lys Glu Glu
             1125 1130
Ala Arg Lys Phe Phe Ala Gly Ile Tyr Glu Glu Ala Phe Asn Gly Ser
                                1150
          1140
                         1145
Trp Ser Thr Lys Thr Asn Trp Leu Phe His Ala Val Lys His Tyr
                       1160
<210> 49
<211> 23
<212> DNA
<213> Artificial Sequence
<220>
<223> Probe/primer derived from C. elegans
```

<210> 50 <211> 20

<400> 49

ggaaatattt taggccagat gcg

<211> 20 <212> DNA 23

```
<213> Artificial Sequence
<220>
<223> Probe/primer derived from C. elegans
<400> 50
                                                                  20
cggacagtcc tgaatacacc
<210> 51
<211> 28
<212> DNA
<213> Artificial Sequence
<223> Probe/primer derived from C. elegans
<400> 51
                                                                  28
tctcqttqtt tgccgtcgga tgtctgcc
<210> 52
<211> 3017
<212> DNA
<213> Caenorhabditis elegans
<400> 52
gtaatcaaat tgtaaaggaa aaatattaat agtcagagta cacataaatg ggtgatcatc 60
ataatttaac gggccttccc ggtacctcca tcccgccaca gttcaactat tctcagcccg 120
gtaccagcac cggaggcccg ctttatggtg gaaaaccttc tcatggattg gaagatattc 180
ctgatgtaga ggaatatgag aggaacctgc tcggggctgg agcaggtttt aatctgctca 240
atgtaggaaa tatggctaat gttcccgacg agcacacacc gatgatgtca ccagtgaata 300
caactacaaa gattctacaa cggagtggta ttaaaatgga aatcccgcca tatttggatc 360
cagacagtca ggatgatgac ccggaagatg gtgtcaacta cccggatcca gatttatttg 420
acacaaaaaa cacaaatatg accgagtacg atttggatgt gttgaagctt ggaaaaccag 480
cagtagatga agcacggaaa aagatcgaag ttcccgacgc tagtgcgccg ccaaacaaaa 540
ttgtagaata tttgatgtat tatagaacgt taaaagaaag tgaactcata caactgaatg 600
cgtatcggac aaaacgaaat cgattatcgt tgaacttggt caaaaacaat attgatcgag 660
agttcgacca aaaagcttgc gagtccctgg tgaaaaaatt gaaggataag aagaatgatc 720
tccagaacct gattgatgtg gttctttcaa aaggtacaaa atataccggt tgcattacaa 780
ttccaaggac acttgatggc cggttacagg tccacggaag aaaaggtttc cctcacgtag 840
tctatggcaa actgtggagg tttaatgaaa tgacaaaaaa cgaaacgcgt catgtggacc 900
actgcaagca cgcatttgaa atgaaaagtg acatggtatg cgtgaatccc tatcactacg 960
aaattgtcat tggaactatg attgttgggc agagggatca tgacaatcga gatatgccgc 1020
cgccacatca acgctaccac actccaggtc ggcaggatcc agttgacgat atgagtagat 1080
ttataccacc agettecatt egteegeete egatgaacat geacacaagg ceteageeta 1140
tgcctcaaca attgccttca gttggcgcaa cgtttgccca tcctctccca catcaggcgc 1200
cacataaccc aggggtttca catccgtact ccattgctcc acagacccat tacccgttga 1260
acatgaaccc aattccgcaa atgccgcaaa tgccacaaat gccaccacct ctccatcagg 1320
gatatggaat gaatgggccg agttgctctt cagaaaacaa caatccattc caccaaaatc 1380
accattataa tgatattagc catccaaatc actattccta cgactgtggt ccgaacttgt 1440
acgggtttcc aactccttat ccggattttc accatccttt caatcagcaa ccacaccagc 1500
cgccacaact atcacaaaac catacgtccc aacaaggcag tcatcaacca gggcaccaag 1560
gtcaggtacc gaatgatcca ccaatttcaa gaccagtgtt acaaccatca acagtcacct 1620
tggacgtgtt ccgtcggtac tgtagacaga catttggaaa tcgatttttt gaaggagaaa 1680
 gtgaacaatc cggcgcaata attcggtcta gtaacaaatt cattgaagaa tttgattcgc 1740
cgatttgtgg tgtgacagtt gttcgaccgc ggatgacaga cggtgaggtt ttggagaaca 1800
 tcatgccgga agatgcacca tatcatgaca tttgcaagtt cattttgagg ctcacatcag 1860
 aaagtgtaac tttctcagga gaggggccag aagttagtga tttgaacgaa aaatggggaa 1920
 caattgtgta ctatgagaaa aatttgcaaa ttggcgagaa aaaatgttcg agaggaaatt 1980
```

```
tccacgtgga tggcggattc atttgctctg agaatcgtta cagtctcgga cttgagccaa 2040
atccaattag agaaccagtg gcgtttaaag ttcgtaaagc aatagtggat ggaattcgct 2100
tttcctacaa aaaagacggg agtgtttggc ttcaaaaccg catgaagtac ccggtatttg 2160
tcacttctgg gtatctcgac gagcaatcag gaggcctaaa gaaggataaa gtgcacaaag 2220
tttacggatg tgcgtctatc aaaacgtttg gcttcaacgt ttccaaacaa atcatcagag 2280
acgcgcttct ttccaagcaa atggcaacaa tgtacttgca aggaaaattg actccgatga 2340
attatatcta cgagaagaag actcaggaag agctgcgaag ggaagcaaca cgcaccactg 2400
attcattggc caagtactgt tgtgtccgtg tctcgttctg caaaggattt ggagaagcat 2460
acccagaacg cccgtcaatt catgattgtc cagtttggat tgagttgaaa atcaacattg 2520
cctacgattt catggattca atctgccagt acataaccaa ctgcttcgag ccgctaggaa 2580
tggaagattt tgcaaaattg ggaatcaacg tcagtgatga ctaaatgata acttttttca 2640
ctcaccctac tagatactga tttagtctta ttccaaatca tccaacgata tcaaactttt 2700
teetttgaac tttgcatact atgttateac aagtteeaag cagttteaat acaaacatag 2760
gatatgttaa caacttttga taagaatcaa gttaccaact gttcattgtg agctttgagc 2820
tgtatagaag gacaatgtat cccatacctc aatctttaat agtcatcagt cactggtccc 2880
gcaccaattt tttcgattcg catatgtcat atattgcacc gtggcccttt ttattgtaac 2940
ttttaatata ttttcttccc aacttgtgaa tatgattgat gaaccaccat tttgagtaat 3000
                                                                  3017
aaatgtattt tttgtgg
<210> 53
<211> 3119
<212> DNA
<213> Caenorhabditis elegans
<400> 53
gtaatcaaat tgtaaaggaa aaatattaat agtcagagta cacataaatg ggtgatcatc 60
ataatttaac gggccttccc ggtacctcca tcccgccaca gttcaactat tctcagcccg 120
gtaccagcac cggaggcccg ctttatggtg gaaaaccttc tcatggattg gaagatattc 180
ctgatgtaga ggaatatgag aggaacctgc tcggggctgg agcaggtttt aatctgctca 240
atgtaggaaa tatggctaat gaatttaaac caataatcac attggacacg aaaccacctc 300
gtgatgccaa caagtcattg gcattcaatg gcgggttgaa gctaatcact ccgaaaactg 360
aagttcccga cgagcacaca ccgatgatgt caccagtgaa tacaactaca aagattctac 420
aacggagtgg tattaaaatg gaaatcccgc catatttgga tccagacagt caggatgatg 480
acccggaaga tggtgtcaac tacccggatc cagatttatt tgacacaaaa aacacaaata 540
tgaccgagta cgatttggat gtgttgaagc ttggaaaacc agcagtagat gaagcacgga 600
aaaagatcga agttcccgac gctagtgcgc cgccaaacaa aattgtagaa tatttgatgt 660
attatagaac gttaaaagaa agtgaactca tacaactgaa tgcgtatcgg acaaaacgaa 720
atcgattatc gttgaacttg gtcaaaaaca atattgatcg agagttcgac caaaaagctt 780
gcgagtccct ggtgaaaaaa ttgaaggata agaagaatga tctccagaac ctgattgatg 840
tggttctttc aaaaggtaca aaatataccg gttgcattac aattccaagg acacttgatg 900
gccggttaca ggtccacgga agaaaaggtt tccctcacgt agtctatggc aaactgtgga 960
ggtttaatga aatgacaaaa aacgaaacgc gtcatgtgga ccactgcaag cacgcatttg 1020
aaatgaaaag tgacatggta tgcgtgaatc cctatcacta cgaaattgtc attggaacta 1080
tgattgttgg gcagagggat catgacaatc gagatatgcc gccgccacat caacgctacc 1140
acactccagg tcggcaggat ccagttgacg atatgagtag atttatacca ccagcttcca 1200
```

ttegteegee teegatgaac atgeacacaa ggeeteagee tatgeeteaa caattgeett 1260 cagttggege aacgtttgee cateetee cacateagge geeacataac eeaggggttt 1320 cacateegta etecattget eeacagaeee attaceegtt gaacatgaac eeaatteege 1380 aaatgeegea aatgeeacaa atgeeacaca etecatea gggatatgga atgaatggge 1440 cgagttgete tteagaaaac aacaateeat teeaceaaaa teaceattat aatgatatta 1500 geeateeaaa teacetattee tacgaetgtg gteegaactt gtacgggttt eeaacteett 1560 ateeggattt teaceateet tteaateage aaceacacaa geegeeacaa etaceaaaa 1620 accaattee aagaceagge ttacaaceat eaacagteac ettggaeggt teegaactt teegaeggt 1740 actgagaea gaeatttgga aategattt ttgaaggaga aagtgaacaa teeggegeaa 1800 taatteeggee geggatgaea gaeggtgagg tttttggagaa cateatgeeg gaagatgeae 1920 catateatga catttgeaag teeattttga ggeteacate agaaagtgta acttteteag 1920 catateatga catttgeaag tteattttga ggeteacate agaaagtgta acttteteag 1980

```
gagaggggcc agaagttagt gatttgaacg aaaaatgggg aacaattgtg tactatgaga 2040
aaaatttgca aattggcgag aaaaaatgtt cgagaggaaa tttccacgtg gatggcggat 2100
tcatttgctc tgagaatcgt tacagtctcg gacttgagcc aaatccaatt agagaaccag 2160
tggcgtttaa agttcgtaaa gcaatagtgg atggaattcg cttttcctac aaaaaagacg 2220
ggagtgtttg gcttcaaaac cgcatgaagt acccggtatt tgtcacttct gggtatctcg 2280
acgagcaatc aggaggccta aagaaggata aagtgcacaa agtttacgga tgtgcgtcta 2340
tcaaaacgtt tggcttcaac gtttccaaac aaatcatcag agacgcgctt ctttccaagc 2400
aaatggcaac aatgtacttg caaggaaaat tgactccgat gaattatatc tacgagaaga 2460
agactcagga agagctgcga agggaagcaa cacgcaccac tgattcattg gccaagtact 2520
gttgtgtccg tgtctcgttc tgcaaaggat ttggagaagc atacccagaa cgcccgtcaa 2580
ttcatgattg tccagtttgg attgagttga aaatcaacat tgcctacgat ttcatggatt 2640
caatctgcca gtacataacc aactgcttcg agccgctagg aatggaagat tttgcaaaat 2700
tgggaatcaa cgtcagtgat gactaaatga taactttttt cactcaccct actagatact 2760
gatttagtct tattccaaat catccaacga tatcaaactt tttcctttga actttgcata 2820
ctatgttatc acaagttcca agcagtttca atacaaacat aggatatgtt aacaactttt 2880
gataagaatc aagttaccaa ctgttcattg tgagctttga gctgtataga aggacaatgt 2940
atcccatacc tcaatcttta atagtcatca gtcactggtc ccgcaccaat tttttcgatt 3000
cgcatatgtc atatattgca ccgtggccct ttttattgta acttttaata tattttcttc 3060
ccaacttgtg aatatgattg atgaaccacc attttgagta ataaatgtat tttttgtgg 3119
<210> 54
<211> 103
<212> PRT
<213> Caenorhabditis elegans
<400> 54
Lys Lys Thr Thr Thr Arg Arg Asn Ala Trp Gly Asn Met Ser Tyr Ala
                                     10
                 5
 1
Glu Leu Ile Thr Thr Ala Ile Met Ala Ser Pro Glu Lys Arg Leu Thr
                                                     30
                                 25
Leu Ala Gln Val Tyr Glu Trp Met Val Gln Asn Val Pro Tyr Phe Arg
Asp Lys Gly Asp Ser Asn Ser Ser Ala Gly Trp Lys Asn Ser Ile Arg
                                             60
    50
                        55
His Asn Leu Ser Leu His Ser Arg Phe Met Arg Ile Gln Asn Glu Gly
                                        75
                    70
Ala Gly Lys Ser Ser Trp Trp Val Ile Asn Pro Asp Ala Lys Pro Gly
                                                         95
Met Asn Pro Arg Arg Thr Arg
            100
<210> 55
<211> 41
<212> PRT
<213> Caenorhabditis elegans
<400> 55
Thr Phe Met Asn Thr Pro Asp Asp Val Met Met Asn Asp Asp Met Glu
                                     10
                  5
 1
Pro Ile Pro Arg Asp Arg Cys Asn Thr Trp Pro Met Arg Arg Pro Gln
                                                     30
                                 25
Leu Glu Pro Pro Leu Asn Ser Ser Pro
```

<210> 56 <211> 109

<212> PRT <213> Caenorhabditis elegans

 Asp Asp Thr Val
 Ser Gly
 Lys
 Lys
 Thr Thr Thr Arg
 Arg
 Asn Ala
 Trp

 1
 5
 10
 15
 15
 15

 Gly
 Asn Met
 Ser Tyr
 Ala
 Glu
 Leu
 Ile
 Thr Thr Thr Ala
 Ile
 Met
 Ala
 Ser

 Pro
 Glu
 Lys
 Arg
 Leu
 Thr Leu
 Ala
 Gln
 Val
 Tyr
 Glu
 Trp
 Met
 Val
 Gln

 Asn
 Val
 Pro
 Tyr
 Phe
 Arg
 Asp
 Lys
 Gly
 Asp
 Ser
 Asn
 Ser
 Ser
 Ala
 Gly

 Trp
 Lys
 Asn
 Ser
 His
 Asn
 Leu
 Ser
 Leu
 His
 Ser
 Arg
 Phe
 Met

 Asp
 Ile
 Glu
 Gly
 Ala
 Gly
 Lys
 Ser
 Ser
 Trp
 Trp
 Val
 Ile
 Asn

 Asp
 Ile
 Glu
 Gly
 Ala
 Gly
 Lys
 Ser
 Ser
 Trp
 Trp
 Val

<210> 57 <211> 655 <212> PRT <213> Homo sapiens

<400> 57 Met Ala Glu Ala Pro Gln Val Val Glu Ile Asp Pro Asp Phe Glu Pro 10 Leu Pro Arg Pro Arg Ser Cys Thr Trp Pro Leu Pro Arg Pro Glu Phe 25 Ser Gln Ser Asn Ser Ala Thr Ser Ser Pro Ala Pro Ser Gly Ser Ala Ala Ala Asn Pro Asp Ala Ala Ala Gly Leu Pro Ser Ala Ser Ala Ala 60 55 Ala Val Ser Ala Asp Phe Met Ser Asn Leu Ser Leu Leu Glu Glu Ser 75 70 Glu Asp Phe Pro Gln Ala Pro Gly Ser Val Ala Ala Ala Val Ala Ala 90 85 Ala Ala Ala Ala Ala Thr Gly Gly Leu Cys Gly Asp Phe Gln Gly 105 Pro Glu Ala Gly Cys Leu His Pro Ala Pro Pro Gln Pro Pro Pro 120 Gly Pro Val Ser Gln His Pro Pro Val Pro Pro Ala Ala Gly Pro 135 140 Leu Ala Gly Gln Pro Arg Lys Ser Ser Ser Ser Arg Arg Asn Ala Trp 155 150 Gly Asn Leu Ser Tyr Ala Asp Leu Ile Thr Lys Ala Ile Glu Ser Ser 170 165 Ala Glu Lys Arg Leu Thr Leu Ser Gln Ile Tyr Glu Trp Met Val Lys 185 Ser Val Pro Tyr Phe Lys Asp Lys Gly Asp Ser Asn Ser Ser Ala Gly 205 200 Trp Lys Asn Ser Ile Arg His Asn Leu Ser Leu His Ser Lys Phe Ile 215 Arg Val Gln Asn Glu Gly Thr Gly Lys Ser Ser Trp Trp Met Leu Asn 235 Pro Glu Gly Gly Lys Ser Gly Lys Ser Pro Arg Arg Ala Ala Ser

```
Met Asp Asn Asn Ser Lys Phe Ala Lys Ser Arg Ser Arg Ala Ala Lys
                              265
Lys Lys Ala Ser Leu Gln Ser Gly Gln Glu Gly Ala Gly Asp Ser Pro
                                              285
                          280
Gly Ser Gln Phe Ser Lys Trp Pro Ala Ser Pro Gly Ser His Ser Asn
                                          300
                      295
Asp Asp Phe Asp Asn Trp Ser Thr Phe Arg Pro Arg Thr Ser Ser Asn
                                      315
               310
Ala Ser Thr Ile Ser Gly Arg Leu Ser Pro Ile Met Thr Glu Gln Asp
                                  330
               325
Asp Leu Gly Glu Gly Asp Val His Ser Met Val Tyr Pro Pro Ser Ala
                              345
Ala Lys Met Ala Ser Thr Leu Pro Ser Leu Ser Glu Ile Ser Asn Pro
                          360
       355
Glu Asn Met Glu Asn Leu Leu Asp Asn Leu Asn Leu Leu Ser Ser Pro
                       375
                                         380
Thr Ser Leu Thr Val Ser Thr Gln Ser Ser Pro Gly Thr Met Met Gln
                  390
                                      395
Gln Thr Pro Cys Tyr Ser Phe Ala Pro Pro Asn Thr Ser Leu Asn Ser
                                  410
               405
Pro Ser Pro Asn Tyr Gln Lys Tyr Thr Tyr Gly Gln Ser Ser Met Ser
                              425
           420
Pro Leu Pro Gln Met Pro Ile Gln Thr Leu Gln Asp Asn Lys Ser Ser
                          440
Tyr Gly Gly Met Ser Gln Tyr Asn Cys Ala Pro Gly Leu Leu Lys Glu
                                          460
            455
Leu Leu Thr Ser Asp Ser Pro Pro His Asn Asp Ile Met Thr Pro Val
                           475
                   470
Asp Pro Gly Val Ala Gln Pro Asn Ser Arg Val Leu Gly Gln Asn Val
                                  490
               485
Met Met Gly Pro Asn Ser Val Met Ser Thr Tyr Gly Ser Gln Ala Ser
                                                  510
                              505
His Asn Lys Met Met Asn Pro Ser Ser His Thr His Pro Gly His Ala
                                              525
                          520
Gln Gln Thr Ser Ala Val Asn Gly Arg Pro Leu Pro His Thr Val Ser
                       535
                                540
Thr Met Pro His Thr Ser Gly Met Asn Arg Leu Thr Gln Val Lys Thr
                                     555
                   550
Pro Val Gln Val Pro Leu Pro His Pro Met Gln Met Ser Ala Leu Gly
                                   570
               565
Gly Tyr Ser Ser Val Ser Ser Cys Asn Gly Tyr Gly Arg Met Gly Leu
                               585
            580
Leu His Gln Glu Lys Leu Pro Ser Asp Leu Asp Gly Met Phe Ile Glu
                                             605
                           600
Arg Leu Asp Cys Asp Met Glu Ser Ile Ile Arg Asn Asp Leu Met Asp
                       615
Gly Asp Thr Leu Asp Phe Asn Phe Asp Asn Val Leu Pro Asn Gln Ser
                                      635
                    630
Phe Pro His Ser Val Lys Thr Thr Thr His Ser Trp Val Ser Gly
                                   650
                645
```

<210> 58

<211> 98

<212> PRT

<213> Caenorhabditis elegans

<400> 58

```
Lys Pro Asn Pro Trp Gly Glu Glu Ser Tyr Ser Asp Ile Ile Ala Lys
Ala Leu Glu Ser Ala Pro Asp Gly Arg Leu Lys Leu Asn Glu Ile Tyr
                                2.5
            20
Gln Trp Phe Ser Asp Asn Ile Pro Tyr Phe Gly Glu Arg Ser Ser Pro
                            40
Glu Glu Ala Ala Gly Trp Lys Asn Ser Ile Arg His Asn Leu Ser Leu
                                            60
                        55
His Ser Arg Phe Met Arg Ile Gln Asn Glu Gly Ala Gly Lys Ser Ser
                                        75
                    70
Trp Trp Val Ile Asn Pro Asp Ala Lys Pro Gly Met Asn Pro Arg Arg
                                    90
Thr Arg
<210> 59
<211> 7
<212> PRT
<213> Caenorhabditis elegans
<400> 59
Trp Lys Asn Ser Ile Arg His
1
<210> 60
<211> 121
<212> PRT
<213> Caenorhabditis elegans
<400> 60
Gln Val Leu Asp Asp His Asp Tyr Gly Arg Cys Val Asp Trp Trp Gly
 1
Val Gly Val Val Met Tyr Glu Met Met Cys Gly Arg Leu Pro Phe Tyr
Ser Lys Asp His Asn Lys Leu Phe Glu Leu Ile Met Ala Gly Asp Leu
        35
Arg Phe Pro Ser Lys Leu Ser Gln Glu Ala Arg Thr Leu Leu Thr Gly
                        55
Leu Leu Val Lys Asp Pro Thr Gln Arg Leu Gly Gly Pro Glu Asp
                                         75
Ala Leu Glu Ile Cys Arg Ala Asp Phe Phe Arg Thr Val Asp Trp Glu
                                     90
                85
Ala Thr Tyr Arg Lys Glu Ile Glu Pro Pro Tyr Lys Pro Asn Val Gln
Ser Glu Thr Asp Thr Ser Tyr Phe Asp
<210> 61
<211> 66
<212> PRT
<213> Caenorhabditis elegans
<400> 61
Thr Met Glu Asp Phe Asp Phe Leu Lys Val Leu Gly Lys Gly Thr Phe
```

```
Gly Lys Val Ile Leu Cys Lys Glu Lys Arg Thr Gln Lys Leu Tyr Ala
Ile Lys Ile Leu Lys Lys Asp Val Ile Ile Ala Arg Glu Glu Val Ala
                                                45
                           40
His Thr Leu Thr Glu Asn Arg Val Leu Gln Arg Cys Lys His Pro Phe
                        55
Leu Thr
65
<210> 62
<211> 45
<212> PRT
<213> Caenorhabditis elegans
<400> 62
Lys Leu Glu Asn Leu Leu Leu Asp Lys Asp Gly His Ile Lys Ile Ala
                                    10
                 5
Asp Phe Gly Leu Cys Lys Glu Glu Ile Ser Phe Gly Asp Lys Thr Ser
                                25
            20
Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val
                            40
<210> 63
<211> 57
<212> PRT
<213> Caenorhabditis elegans
<400> 63
Tyr Phe Gln Glu Leu Lys Tyr Ser Phe Gln Glu Gln His Tyr Leu Cys
                 5
1
Phe Val Met Gln Phe Ala Asn Gly Gly Glu Leu Phe Thr His Val Arg
                                 25
Lys Cys Gly Thr Phe Ser Glu Pro Arg Ala Arg Phe Tyr Gly Ala Glu
        35
Ile Val Leu Ala Leu Gly Tyr Leu His
                         55
    50
<210> 64
<211> 59
<212> PRT
<213> Caenorhabditis elegans
<400> 64
Ser Thr Phe Ala Ile Phe Tyr Phe Gln Thr Met Leu Phe Glu Lys Pro
                                                         15
 1
                 5
                                     10
Arg Pro Asn Met Phe Met Val Arg Cys Leu Gln Trp Thr Thr Val Ile
                                 25
Glu Arg Thr Phe Tyr Ala Glu Ser Ala Glu Val Arg Gln Arg Trp Ile
His Ala Ile Glu Ser Ile Ser Lys Lys Tyr Lys
```

<210> 65 <211> 33

```
<212> PRT
<213> Caenorhabditis elegans
<400> 65
Leu Gln Glu Leu Lys Tyr Ser Phe Gln Thr Asn Asp Arg Leu Cys Phe
                                     10
Val Met Glu Phe Ala Ile Gly Gly Asp Leu Tyr Tyr His Leu Asn Arg
                                25
Glu
<210> 66
<211> 21
<212> PRT
<213> Caenorhabditis elegans
<400> 66
Val Val Ile Glu Gly Trp Leu His Lys Lys Gly Glu His Ile Arg Asn
                 5
Trp Arg Pro Arg Phe
            20
<210> 67
<211> 26
<212> PRT
<213> Caenorhabditis elegans
<400> 67
Phe Ser Glu Pro Arg Ala Arg Phe Tyr Gly Ser Glu Ile Val Leu Ala
                 5
Leu Gly Tyr Leu His Ala Asn Ser Ile Val
            20
<210> 68
<211> 39
<212> PRT
<213> Caenorhabditis elegans
<400> 68
Ile Arg Val Ser Phe Cys Lys Gly Phe Gly Glu Thr Tyr Ser Arg Leu
                                     10
                  5
Lys Val Val Asn Leu Pro Cys Trp Ile Glu Ile Ile Leu His Glu Pro
                                 25
Ala Asp Glu Tyr Asp Thr Val
        35
<210> 69
<211> 45
<212> PRT
<213> Caenorhabditis elegans
<400> 69
Ser Arg Asn Ser Lys Ser Ser Gln Ile Arg Asn Thr Val Gly Ala Gly
                                     10
```

```
Ile Gln Leu Ala Tyr Glu Asn Gly Glu Leu Trp Leu Thr Val Leu Thr
                                25
Asp Gln Ile Val Phe Val Gln Cys Pro Phe Leu Asn Gln
<210> 70
<211> 29
<212> PRT
<213> Caenorhabditis elegans
<400> 70
Asn Glu Met Leu Asp Pro Glu Pro Lys Tyr Pro Lys Glu Glu Lys Pro
                5
                                    10
Trp Cys Thr Ile Phe Tyr Tyr Glu Leu Thr Val Arg Val
<210> 71
<211> 29
<212> PRT
<213> Caenorhabditis elegans
<400> 71
Gln Leu Gly Lys Ala Phe Glu Ala Lys Val Pro Thr Ile Thr Ile Asp
                5
                                    1.0
Gly Ala Thr Gly Ala Ser Asp Glu Cys Arg Met Ser Leu
            20
<210> 72
<211> 105
<212> PRT
<213> Caenorhabditis elegans
<400> 72
Ser Pro Asp Asp Gly Leu Leu Asp Ser Ser Glu Glu Ser Arg Arg
                                    10
Gln Lys Thr Cys Arg Val Cys Gly Asp His Ala Thr Gly Tyr Asn Phe
                                25
Asn Val Ile Thr Cys Glu Ser Cys Lys Ala Phe Phe Arg Arg Asn Ala
                            40
                                                45
Leu Arg Pro Lys Glu Phe Lys Cys Pro Tyr Ser Glu Asp Cys Glu Ile
                        55
Asn Ser Val Ser Arg Arg Phe Cys Gln Lys Cys Arg Leu Arg Lys Cys
                                        75
Phe Thr Val Gly Met Lys Lys Glu Trp Ile Leu Asn Glu Glu Gln Leu
Arg Arg Lys Asn Ser Arg Leu Asn
            100
<210> 73
<211> 89
<212> PRT
<213> Caenorhabditis elegans
<400> 73
```

```
Leu Asp Ser Ser Glu Glu Ser Arg Arg Arg Gln Lys Thr Cys Arg Val
                                    10
Cys Gly Asp His Ala Thr Gly Tyr Asn Phe Asn Val Ile Thr Cys Glu
                                25
Ser Cys Lys Ala Phe Phe Arg Arg Asn Ala Leu Arg Pro Lys Glu Phe
Lys Cys Pro Tyr Ser Glu Asp Cys Glu Ile Asn Ser Val Ser Arg Arg
                        55
Phe Cys Gln Lys Cys Arg Leu Arg Lys Cys Phe Thr Val Gly Met Lys
                   70
Lys Glu Trp Ile Leu Asn Glu Glu Gln
                85
```

<210> 74 <211> 73 <212> PRT <213> Caenorhabditis elegans

<400> 74

Asp Ile Met Asn Ile Met Asp Val Thr Met Arg Arg Phe Val Lys Val 1 Ala Lys Gly Val Pro Ala Phe Arg Glu Val Ser Gln Glu Gly Lys Phe 20 25 Ser Leu Leu Lys Gly Gly Met Ile Glu Met Leu Thr Val Arg Gly Val 40 Thr Arg Tyr Asp Ala Ser Thr Asn Ser Phe Lys Thr Pro Thr Ile Lys 55 Gly Gln Asn Val Ser Val Asn Val Asp

<210> 75 <211> 112 <212> PRT <213> Caenorhabditis elegans

<400> 75 Ser Gly Ser Leu Val Asp Leu Met Ile Lys Asn Leu Thr Ala Tyr Thr 10 Gln Gly Leu Asn Glu Thr Val Lys Asn Arg Thr Ala Glu Leu Glu Lys 25 Glu Gln Glu Lys Gly Asp Gln Leu Leu Met Glu Leu Leu Pro Lys Ser 40 Val Ala Asn Asp Leu Lys Asn Gly Ile Ala Val Asp Pro Lys Val Tyr 55 60 Glu Asn Ala Thr Ile Leu Tyr Ser Asp Ile Val Gly Phe Thr Ser Leu 75 Cys Ser Gln Ser Gln Pro Met Glu Val Val Thr Leu Leu Ser Gly Met 90 Tyr Gln Arg Phe Asp Leu Ile Ile Ser Gln Gln Gly Gly Tyr Lys Val

<210> 76 <211> 107 <212> PRT <213> Caenorhabditis elegans

Arg Glu Asp

```
<400> 76
Met Glu Thr Ile Gly Asp Ala Tyr Cys Val Ala Ala Gly Leu Pro Val
                                    10
Val Met Glu Lys Asp His Val Lys Ser Ile Cys Met Ile Ala Leu Leu
Gln Arg Asp Cys Leu His His Phe Glu Ile Pro His Arg Pro Gly Thr
                            40
Phe Leu Asn Cys Arg Trp Gly Phe Asn Ser Gly Pro Val Phe Ala Gly
                        55
Val Ile Gly Gln Lys Ala Pro Arg Tyr Ala Cys Phe Gly Glu Ala Val
                                        75
                    70
Ile Leu Ala Ser Lys Met Glu Ser Ser Gly Val Glu Asp Arg Ile Gln
                                   90
Met Thr Leu Ala Ser Gln Gln Leu Leu Glu Glu
                                105
            100
<210> 77
<211> 43
<212> PRT
<213> Caenorhabditis elegans
<400> 77
Asp Ile Leu Lys Gly Leu Glu Tyr Ile His Ala Ser Ala Ile Asp Phe
                 -5
                                    10
His Gly Asn Leu Thr Leu His Asn Cys Met Leu Asp Ser His Trp Ile
                                25
Val Lys Leu Ser Gly Phe Gly Val Asn Arg Leu
                            40
        35
<210> 78
<211> 15
<212> PRT
<213> Caenorhabditis elegans
<400> 78
Asp Met Tyr Ser Phe Gly Val Ile Leu His Glu Ile Ile Leu Lys
 1
<210> 79
<211> 67
<212> PRT
<213> Caenorhabditis elegans
<400> 79
Ala Ile Lys Ile Asn Val Asp Asp Pro Ala Ser Thr Glu Asn Leu Asn
 1
Tyr Leu Met Glu Ala Asn Ile Met Lys Asn Phe Lys Thr Asn Phe Ile
                                 25
            20
Val Gln Leu Tyr Gly Val Ile Ser Thr Val Gln Pro Ala Met Val Val
                             40
Met Glu Met Met Asp Leu Gly Asn Leu Arg Asp Tyr Leu Arg Ser Lys
                         55
    50
```

```
<210> 80
<211> 54
<212> PRT
<213> Caenorhabditis elegans
Val Ile Lys Lys Pro Glu Cys Cys Glu Asn Tyr Trp Tyr Lys Val Met
                                    10
Lys Met Cys Trp Arg Tyr Ser Pro Arg Asp Arg Pro Thr Phe Leu Gln
                                                     30
            20
                                25
Leu Val His Leu Leu Ala Ala Glu Ala Ser Pro Glu Phe Arg Asp Leu
        35
                            40
Ser Phe Val Leu Thr Asp
    50
<210> 81
<211> 69
<212> PRT
<213> Caenorhabditis elegans
<400> 81
Lys Gln Asp Ser Gly Met Ala Ser Glu Leu Lys Asp Ile Phe Ala Asn
                                    10
1
Ile His Thr Ile Thr Gly Tyr Leu Leu Val Arg Gln Ser Ser Pro Phe
                                25
Ile Ser Leu Asn Met Phe Arg Asn Leu Arg Arg Ile Glu Ala Lys Ser
                            40
                                                 45
Leu Phe Arg Asn Leu Tyr Ala Ile Thr Val Phe Glu Asn Pro Asn Leu
                                             60
Lys Lys Leu Phe Asp
65
<210> 82
<211> 52
<212> PRT
<213> Caenorhabditis elegans
<400> 82
Phe Pro His Leu Arg Glu Ile Thr Gly Thr Leu Leu Val Phe Glu Thr
                                     10
 1
Glu Gly Leu Val Asp Leu Arg Lys Ile Phe Pro Asn Leu Arg Val Ile
                                 25
Gly Gly Arg Ser Leu Ile Gln His Tyr Ala Leu Ile Ile Tyr Arg Asn
        35
Pro Asp Leu Glu
    50
<210> 83
<211> 46
<212> PRT
<213> Caenorhabditis elegans
<400> 83
Glu Ile Gly Leu Asp Lys Leu Ser Val Ile Arg Asn Gly Gly Val Arg
                  5
```

```
Ile Ile Asp Asn Arg Lys Leu Cys Tyr Thr Lys Thr Ile Asp Trp Lys
His Leu Ile Thr Ser Ser Ile Asn Asp Val Val Val Asp Asn
        35
                            40
<210> 84
<211> 36
<212> PRT
<213> Caenorhabditis elegans
<400> 84
Tyr Asn Ala Asp Asp Trp Glu Leu Arg Gln Asp Asp Val Val Leu Gly
                                     10
Gln Gln Cys Gly Glu Gly Ser Phe Gly Lys Val Tyr Leu Gly Thr Gly
            20
Asn Asn Val Val
        35
<210> 85
<211> 24
<212> PRT
<213> Caenorhabditis elegans
<400> 85
Asp Ser Leu Ala Lys Tyr Cys Cys Val Arg Val Ser Phe Cys Lys Gly
                                     10
                 5
Phe Gly Glu Ala Tyr Pro Glu Arg
<210> 86
<211> 13
<212> PRT
<213> Caenorhabditis elegans
<400> 86
Gly Trp Asp Trp Ile Val Ala Pro Pro Arg Tyr Asn Ala
                 5
 1
<210> 87
<211> 121
<212> PRT
<213> Homo sapiens
<400> 87
Glu Val Leu Glu Asp Asn Asp Tyr Gly Arg Ala Val Asp Trp Trp Gly
                                     10
                 5
Leu Gly Val Val Met Tyr Glu Met Met Cys Gly Arg Leu Pro Phe Tyr
                                 25
Asn Gln Asp His Glu Lys Leu Phe Glu Leu Ile Leu Met Glu Glu Ile
                                                 45
Arg Phe Pro Arg Thr Leu Gly Pro Glu Ala Lys Ser Leu Leu Ser Gly
                         55
Leu Leu Lys Lys Asp Pro Thr Gln Arg Leu Gly Gly Ser Glu Asp
                                                              80
```

```
Ala Lys Glu Ile Met Gln His Arg Phe Phe Ala Asn Ile Val Trp Gln
                                    90
Asp Val Tyr Glu Lys Lys Leu Ser Pro Pro Phe Lys Pro Gln Val Thr
                               105
            100
Ser Glu Thr Asp Thr Arg Tyr Phe Asp
                            120
        115
<210> 88
<211> 121
<212> PRT
<213> Caenorhabditis elegans
<400> 88
Gln Val Leu Asp Asp His Asp Tyr Gly Arg Cys Val Asp Trp Trp Gly
Val Gly Val Val Met Tyr Glu Met Met Cys Gly Arg Leu Pro Phe Tyr
                                25
            20
Ser Lys Asp His Asn Lys Leu Phe Glu Leu Ile Met Ala Gly Asp Leu
                            40
Arg Phe Pro Ser Lys Leu Ser Gln Glu Ala Arg Thr Leu Leu Thr Gly
                        55
Leu Leu Val Lys Asp Pro Thr Gln Arg Leu Gly Gly Pro Glu Asp
                    70
                                        75
Ala Leu Glu Ile Cys Arg Ala Asp Phe Phe Arg Thr Val Asp Trp Glu
                                   90
                85
Ala Thr Tyr Arg Lys Glu Ile Glu Pro Pro Tyr Lys Pro Asn Val Gln
                                105
Ser Glu Thr Asp Thr Ser Tyr Phe Asp
                            120
<210> 89
<211> 66
<212> PRT
<213> Homo sapiens
<400> 89
Thr Met Asn Glu Phe Glu Tyr Leu Lys Leu Leu Gly Lys Gly Thr Phe
                                    10
 1
Gly Lys Val Ile Leu Val Lys Glu Lys Ala Thr Gly Arg Tyr Tyr Ala
                                                     30
                                25
Met Lys Ile Leu Lys Lys Glu Val Ile Val Ala Lys Asp Glu Val Ala
                                                45
His Thr Leu Thr Glu Asn Arg Val Leu Gln Asn Ser Arg His Pro Phe
    50
Leu Thr
65
<210> 90
 <211> 66
 <212> PRT
 <213> Caenorhabditis elegans
 <400> 90
 Thr Met Glu Asp Phe Asp Phe Leu Lys Val Leu Gly Lys Gly Thr Phe
```

```
Gly Lys Val Ile Leu Cys Lys Glu Lys Arg Thr Gln Lys Leu Tyr Ala
Ile Lys Ile Leu Lys Lys Asp Val Ile Ile Ala Arg Glu Glu Val Ala
                                                 45
                           40
His Thr Leu Thr Glu Asn Arg Val Leu Gln Arg Cys Lys His Pro Phe
   50
Leu Thr
65
<210> 91
<211> 45
<212> PRT
<213> Homo sapiens
<400> 91
Lys Leu Glu Asn Leu Met Leu Asp Lys Asp Gly His Ile Lys Ile Thr
                                     10
                 5
Asp Phe Gly Leu Cys Lys Glu Gly Ile Lys Asp Gly Ala Thr Met Lys
                                 25
            2.0
Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val
        35
<210> 92
<211> 45
<212> PRT
<213> Caenorhabditis elegans
<400> 92
Lys Leu Glu Asn Leu Leu Leu Asp Lys Asp Gly His Ile Lys Ile Ala
                                     10
 1
                 5
Asp Phe Gly Leu Cys Lys Glu Glu Ile Ser Phe Gly Asp Lys Thr Ser
                                25
Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val
                             40
<210> 93
<211> 57
<212> PRT
<213> Homo sapiens
<400> 93
Phe Leu Thr Ala Leu Lys Tyr Ser Phe Gln Thr His Asp Arg Leu Cys
                                     10
 1
 Phe Val Met Glu Tyr Ala Asn Gly Gly Glu Leu Phe Phe His Leu Ser
                                                     3.0
             20
                                 25
 Arg Glu Arg Val Phe Ser Glu Asp Arg Ala Arg Phe Tyr Gly Ala Glu
                                                  45
 Ile Val Ser Ala Leu Asp Tyr Leu His
                         55
     50
 <210> 94
 <211> 57
 <212> PRT
 <213> Caenorhabditis elegans
```

```
<400> 94
Tyr Phe Gln Glu Leu Lys Tyr Ser Phe Gln Glu Gln His Tyr Leu Cys
Phe Val Met Gln Phe Ala Asn Gly Gly Glu Leu Phe Thr His Val Arg
                                25
Lys Cys Gly Thr Phe Ser Glu Pro Arg Ala Arg Phe Tyr Gly Ala Glu
                           40
       35
Ile Val Leu Ala Leu Gly Tyr Leu His
<210> 95
```

<211> 59 <212> PRT

<213> Homo sapiens

<400> 95

Asn Asn Phe Ser Val Ala Gln Cys Gln Leu Met Lys Thr Glu Arg Pro 5 Arg Pro Asn Thr Phe Ile Ile Arg Cys Leu Gln Trp Thr Thr Val Ile 30 20 Glu Arg Thr Phe His Val Glu Thr Pro Glu Glu Arg Glu Glu Trp Ala 40

Thr Ala Ile Gln Thr Val Ala Asp Gly Leu Lys 55

<210> 96 <211> 59 <212> PRT <213> Caenorhabditis elegans

<400> 96 Ser Thr Phe Ala Ile Phe Tyr Phe Gln Thr Met Leu Phe Glu Lys Pro 10 5 Arg Pro Asn Met Phe Met Val Arg Cys Leu Gln Trp Thr Thr Val Ile 30 25 Glu Arg Thr Phe Tyr Ala Glu Ser Ala Glu Val Arg Gln Arg Trp Ile 40 His Ala Ile Glu Ser Ile Ser Lys Lys Tyr Lys 50

<210> 97 <211> 33 <212> PRT <213> Homo sapiens

<400> 97 Leu Thr Ala Leu Lys Tyr Ser Phe Gln Thr His Asp Arg Leu Cys Phe 10 Val Met Glu Tyr Ala Asn Gly Gly Glu Leu Phe Phe His Leu Ser Arg 20 25 30 Glu

<210> 98

```
<211> 33
<212> PRT
<213> Caenorhabditis elegans
<400> 98
Leu Gln Glu Leu Lys Tyr Ser Phe Gln Thr Asn Asp Arg Leu Cys Phe
                                    10
Val Met Glu Phe Ala Ile Gly Gly Asp Leu Tyr Tyr His Leu Asn Arg
                                25
Glu
<210> 99
<211> 473
<212> PRT
<213> Homo sapiens
<400> 99
Met Leu Gly Thr Val Lys Met Glu Gly His Glu Thr Ser Asp Trp Asn
                                    10
                 5
Ser Tyr Tyr Ala Asp Thr Gln Glu Ala Tyr Ser Ser Val Pro Val Ser
                                25
Asn Met Asn Ser Gly Leu Gly Ser Met Asn Ser Met Asn Thr Tyr Met
                                                45
                            40
Thr Met Asn Thr Met Thr Thr Ser Gly Asn Met Thr Pro Ala Ser Phe
                        55
                                            60
Asn Met Ser Tyr Ala Asn Pro Ala Leu Gly Ala Gly Leu Ser Pro Gly
                                        75
                    70
Ala Val Ala Gly Met Pro Gly Gly Ser Ala Gly Ala Met Asn Ser Met
                                    90
Thr Ala Ala Gly Val Thr Ala Met Gly Thr Ala Leu Ser Pro Ser Gly
                                                    110
                                105
Met Gly Ala Met Gly Ala Gln Gln Ala Ala Ser Met Met Asn Gly Leu
                                                125
                            120
Gly Pro Tyr Ala Ala Ala Met Asn Pro Cys Met Ser Pro Met Ala Tyr
                                            140
                        135
Ala Pro Ser Asn Leu Gly Arg Ser Arg Ala Gly Gly Gly Asp Ala
                                        155
                    150
Lys Thr Phe Lys Arg Ser Tyr Pro His Ala Lys Pro Pro Tyr Ser Tyr
                                    170
                165
Ile Ser Leu Ile Thr Met Ala Ile Gln Arg Ala Pro Ser Lys Met Leu
                                185
                                                     190
Thr Leu Ser Glu Ile Tyr Gln Trp Ile Met Asp Leu Phe Pro Tyr Tyr
                                                 205
                            200
Arg Gln Asn Gln Gln Arg Trp Gln Asn Ser Ile Arg His Ser Leu Ser
                                             220
                         215
Phe Asn Asp Cys Phe Val Lys Val Ala Arg Ser Pro Asp Lys Pro Gly
                                         235
                     230
Lys Gly Ser Tyr Trp Thr Leu His Pro Asp Ser Gly Asn Met Phe Glu
                                     250
                 245
Asn Gly Cys Tyr Leu Arg Arg Gln Lys Arg Phe Lys Cys Glu Lys Gln
                                 265
            260
 Pro Gly Ala Gly Gly Gly Gly Ser Gly Ser Gly Gly Ser Gly Ala
                                                 285
                             280
Lys Gly Gly Pro Glu Ser Arg Lys Asp Pro Ser Gly Ala Ser Asn Pro
                         295
```

Ser Ala Asp Ser Pro Leu His Arg Gly Val His Gly Lys Thr Gly Gln

```
315
                   310
305
Leu Glu Gly Ala Pro Ala Pro Gly Pro Ala Ala Ser Pro Gln Thr Leu
                                   330
               325
Asp His Ser Gly Ala Thr Ala Thr Gly Gly Ala Ser Glu Leu Lys Thr
                               345
           340
Pro Ala Ser Ser Thr Ala Pro Pro Ile Ser Ser Gly Pro Gly Ala Leu
                                               365
                           360
Ala Ser Val Pro Ala Ser His Pro Ala His Gly Leu Ala Pro His Glu
                                           380
                       375
Ser Gln Leu His Leu Lys Gly Asp Pro His Tyr Ser Phe Asn His Pro
                   390
                                       395
Phe Ser Ile Asn Asn Leu Met Ser Ser Ser Glu Gln Gln His Lys Leu
                                   410
               405
Asp Phe Lys Ala Tyr Glu Gln Ala Leu Gln Tyr Ser Pro Tyr Gly Ser
                               425
                                                   430
Thr Leu Pro Ala Ser Leu Pro Leu Gly Ser Ala Ser Val Thr Thr Arg
                        440
Ser Pro Ile Glu Pro Ser Ala Leu Glu Pro Ala Tyr Tyr Gln Gly Val
                    455
Tyr Ser Arg Pro Val Leu Asn Thr Ser
                    470
<210> 100
<211> 347
<212> PRT
<213> Homo sapiens
<400> 100
Met Leu Gly Ser Val Lys Met Glu Ala His Asp Leu Ala Glu Trp Ser
                                    10
```

Tyr Tyr Pro Glu Ala Gly Glu Val Tyr Ser Pro Val Thr Pro Val Pro 25 Thr Met Ala Pro Leu Asn Ser Tyr Met Thr Leu Asn Pro Leu Ser Ser 40 Pro Tyr Pro Gly Gly Leu Pro Ala Ser Pro Leu Pro Ser Gly Pro Leu 55 60 Ala Pro Pro Ala Pro Ala Ala Pro Leu Gly Pro Thr Phe Pro Gly Leu 75 70 Gly Leu Ser Gly Gly Ser Ser Ser Gly Tyr Gly Ala Pro Gly Pro 90 85 Gly Leu Val His Gly Lys Glu Met Pro Lys Gly Tyr Arg Ala Pro Ala 105 100 His Ala Lys Pro Pro Tyr Ser Tyr Ile Ser Leu Ile Thr Met Ala Ile 115 120 Gln Gln Ala Pro Gly Lys Val Leu Thr Leu Ser Glu Ile Tyr Gln Trp 140 135 Ile Met Asp Leu Phe Pro Tyr Tyr Arg Asp Asn Gln Gln Arg Trp Gln 155 150 Asn Ser Ile Arg His Ser Leu Ser Phe Asn Asp Cys Phe Val Lys Val 170 165 Ala Arg Ser Pro Asp Lys Pro Gly Lys Gly Ser Tyr Trp Ala Leu His 185 180 Pro Ser Ser Gly Asn Met Phe Glu Asn Gly Cys Tyr Leu Arg Arg Gln 200 205 195 Lys Arg Phe Lys Leu Glu Glu Lys Val Lys Lys Gly Gly Ser Gly Ala Ser Thr Thr Arg Asn Gly Thr Gly Ser Ala Ala Ser Thr Thr Thr Pro

```
235
                  230
225
Ala Ala Thr Val Thr Ser Pro Pro Gln Pro Pro Pro Pro Ala Pro Glu
                                 250
               245
Pro Glu Ala Gln Gly Gly Glu Asp Val Gly Ala Leu Asp Cys Gly Ser
                              265
Pro Ala Ser Ser Thr Pro Tyr Phe Thr Gly Leu Glu Leu Pro Gly Asp
                                              285
                           280
Leu Lys Leu Asp Ala Pro Tyr Asn Phe Asn His Pro Phe Ser Ile Asn
                       295
                                          300
Asn Leu Met Ser Glu Gln Thr Pro Ala Pro Pro Lys Leu Asp Val Gly
                   310
                                      315
Phe Gly Gly Tyr Gly Ala Glu Gly Gly Glu Pro Gly Val Tyr Tyr Gln
                        330
               325
Gly Leu Tyr Ser Arg Ser Leu Leu Asn Ala Ser
```

<211> 635

<212> PRT

<213> Caenorhabditis elegans

 $<\!400\!> 101$ Met Met Leu Val Asp Gln Gly Thr Asp Ala Ser Ser Ser Ala

Ser Thr Ser Thr Ser Ser Val Ser Arg Phe Gly Ala Asp Thr Phe Met Asn Thr Pro Asp Asp Val Met Met Asn Asp Asp Met Glu Pro Ile Pro Arg Asp Arg Cys Asn Thr Trp Pro Met Arg Arg Pro Gln Leu Glu Pro Pro Leu Asn Ser Ser Pro Ile Ile His Glu Gln Ile Pro Glu Glu Asp Ala Asp Leu Tyr Gly Ser Asn Glu Gln Cys Gly Gln Leu Gly Gly Ala Ser Ser Asn Gly Ser Thr Ala Met Leu His Thr Pro Asp Gly Ser Asn Ser His Gln Thr Ser Phe Pro Ser Glu Cys Tyr Thr Trp Pro Met Gln Gln Tyr Ile Tyr Gln Glu Ser Ser Ala Thr Ile Pro His His Leu Asn Gln His Asn Asn Pro Tyr His Pro Met His Pro His His Gln Leu Pro His Met Gln Gln Leu Pro Gln Pro Leu Leu Asn Leu Asn Met Thr Thr Leu Thr Ser Ser Gly Ser Ser Val Ala Ser Ser Ile Gly Gly Ala Gln Cys Ser Pro Cys Ala Ser Gly Ser Ser Thr Ala Ala Thr Asn Ser Ser Gln Gln Gln Thr Val Gly Gln Met Leu Ala Ala Ser Val Pro Cys Ser Ser Ser Gly Met Thr Leu Gly Met Ser Leu Asn Leu Ser Gln Gly Gly Gly Pro Met Pro Ala Lys Lys Lys Arg Cys Arg Lys Lys Pro Thr Asp Gln Leu Ala Gln Lys Lys Pro Asn Pro Trp Gly Glu Glu Ser Tyr Ser Asp Ile Ile Ala Lys Ala Leu Glu Ser Ala Pro Asp Gly

```
275
                            280
                                                285
Arg Leu Lys Leu Asn Glu Ile Tyr Gln Trp Phe Ser Asp Asn Ile Pro
                                           300
                       295
Tyr Phe Gly Glu Arg Ser Ser Pro Glu Glu Ala Ala Gly Trp Lys Asn
                                       315
                    310
Ser Ile Arg His Asn Leu Ser Leu His Ser Arg Phe Met Arg Ile Gln
                                    330
                325
Asn Glu Gly Ala Gly Lys Ser Ser Trp Trp Val Ile Asn Pro Asp Ala
                               345
Lys Pro Gly Met Asn Pro Arg Arg Thr Arg Glu Arg Ser Asn Thr Ile
                            360
Glu Thr Thr Thr Lys Ala Gln Leu Glu Lys Ser Arg Arg Gly Ala Lys
                        375
                                            380
Lys Arg Ile Lys Glu Arg Ala Leu Met Gly Ser Leu His Ser Thr Leu
                                        395
                    390
Asn Gly Asn Ser Ile Ala Gly Ser Ile Gln Thr Ile Ser His Asp Leu
                                    410
               405
Tyr Asp Asp Asp Ser Met Gln Gly Ala Phe Asp Asn Val Pro Ser Ser
           420
                               425
Phe Arg Pro Arg Thr Gln Ser Asn Leu Ser Ile Pro Gly Ser Ser Ser
                            440
Arg Val Ser Pro Ala Ile Gly Ser Asp Ile Tyr Asp Asp Leu Glu Phe
                        455
                                            460
Pro Ser Trp Val Gly Glu Ser Val Pro Ala Ile Pro Ser Asp Ile Val
                    470
                                        475
Asp Arg Thr Asp Gln Met Arg Ile Asp Ala Thr Thr His Ile Gly Gly
                                    490
               485
Val Gln Ile Lys Gln Glu Ser Lys Pro Ile Lys Thr Glu Pro Ile Ala
                               505
                                                    510
            500
Pro Pro Pro Ser Tyr His Glu Leu Asn Ser Val Arg Gly Ser Cys Ala
                           520
                                                525
Gln Asn Pro Leu Leu Arg Asn Pro Ile Val Pro Ser Thr Asn Phe Lys
                                            540
                        535
Pro Met Pro Leu Pro Gly Ala Tyr Gly Asn Tyr Gln Asn Gly Gly Ile
                    550
                                       555
Thr Pro Ile Asn Trp Leu Ser Thr Ser Asn Ser Ser Pro Leu Pro Gly
                                    570
                565
Ile Gln Ser Cys Gly Ile Val Ala Ala Gln His Thr Val Ala Ser Ser
                                585
Ser Ala Leu Pro Ile Asp Leu Glu Asn Leu Thr Leu Pro Asp Gln Pro
                            600
Leu Met Asp Thr Met Asp Val Asp Ala Leu Ile Arg His Glu Leu Ser
                        615
Gln Ala Gly Gly Gln His Ile His Phe Asp Leu
                    630
```

```
<210> 102
```

<400> 102

Met Arg Ile Gln Pro Gln Lys Ala Ala Ala Ile Ile Asp Leu Asp Pro 1 5 10 15

Asp Phe Glu Pro Gln Ser Arg Pro Arg Ser Cys Thr Trp Pro Leu Pro 20 25 30

Arg Pro Glu Ile Ala Asn Gln Pro Ser Glu Pro Pro Glu Val Glu Pro

<211> 501

<212> PRT

<213> Homo sapiens

		35					40					45			
-	50					55					60			Ile	
65			-		70					75				Gly	80
				85					90					Asn 95	
Trp	Gly	Asn	Gln 100	Ser	Tyr	Ala	Glu	Phe 105	Ile	Ser	Gln	Ala	Ile 110	Glu	Ser
Ala	Pro	Glu 115	Lys	Arg	Leu	Thr	Leu 120	Ala	Gln	Ile	Tyr	Glu 125	Trp	Met	Val
Arg	Thr 130	Val	Pro	Tyr	Phe	Lys 135	Asp	Lys	Gly	Asp	Ser 140	Asn	Ser	Ser	Ala
Gly 145	Trp	Lys	Asn	Ser	Ile 150	Arg	His	Asn	Leu	Ser 155	Leu	His	Ser	Lys	Phe 160
				165					170					Met 175	
			180					185					190	Ala	
		195					200					205		Ala	
	210					215					220			Thr	
225					230					235				Cys	240
				245					250					Arg 255	
			260					265					270	Arg	
		275					280					285		Ser	
	290	_				295					300			Leu	
305					310					315				Gly	320
	_			325					330					His	
_			340					345					350	Ala	
		355					360					365		Thr	
	370					375					380			Asp	
385					390					395					Ser 400
				405					410					Leu 415	
			420					425					430	Ala	
		435					440					445		Thr	
	450					455					460				Gln
465					470					475				Met	480
				485	ьeu	Met	Asp	Glu	GIY 490	GLU	GТĀ	ьeu	ASP	Phe 495	ASN
Phe	Glu	Pro	Asp	Pro											

```
<210> 103
<211> 366
<212> PRT
<213> Homo sapiens
<400> 103
Arg Gly Ala Ile Arg Ile Glu Lys Asn Ala Asp Leu Cys Tyr Leu Ser
                                   10
Thr Val Asp Trp Ser Leu Ile Leu Asp Ala Val Ser Asn Asn Tyr Ile
                                25
Val Gly Asn Lys Pro Pro Lys Glu Cys Gly Asp Leu Cys Pro Gly Thr
                           40
Met Glu Glu Lys Pro Met Cys Glu Lys Thr Thr Ile Asn Asn Glu Tyr
                        55
Asn Tyr Arg Cys Trp Thr Thr Asn Arg Cys Gln Lys Met Cys Pro Ser
                                        75
                   70
Thr Cys Gly Lys Arg Ala Cys Thr Glu Asn Asn Glu Cys Cys His Pro
                                    90
                85
Glu Cys Leu Gly Ser Cys Ser Ala Pro Asp Asn Asp Thr Ala Cys Val
                                105
                                                    110
Ala Cys Arg His Tyr Tyr Ala Gly Val Cys Val Pro Ala Cys Pro
                           120
                                                125
Pro Asn Thr Tyr Arg Phe Glu Gly Trp Arg Cys Val Asp Arg Asp Phe
                        135
                                           140
Cys Ala Asn Ile Leu Ser Ala Glu Ser Ser Asp Ser Glu Gly Phe Val
                                       155
                    150
Ile His Asp Gly Glu Cys Met Gln Glu Cys Pro Ser Gly Phe Ile Arg
                                    170
Asn Gly Ser Gln Ser Met Tyr Cys Ile Pro Cys Glu Gly Pro Cys Pro
                                185
                                                   190
Lys Val Cys Glu Glu Glu Lys Lys Thr Lys Thr Ile Asp Ser Val Thr
                            200
                                                205
        195
Ser Ala Gln Met Leu Gln Gly Cys Thr Ile Phe Lys Gly Asn Leu Leu
                        215
                                            220
Ile Asn Ile Arg Arg Gly Asn Asn Ile Ala Ser Glu Leu Glu Asn Phe
                    230
                                        235
Met Gly Leu Ile Glu Val Val Thr Gly Tyr Val Lys Ile Arg His Ser
                                    250
                245
His Ala Leu Val Ser Leu Ser Phe Leu Lys Asn Leu Arg Leu Ile Leu
                                265
            260
Gly Glu Glu Gln Leu Glu Gly Asn Tyr Ser Phe Tyr Val Leu Asp Asn
                                                285
                            280
Gln Asn Leu Gln Gln Leu Trp Asp Trp Asp His Arg Asn Leu Thr Ile
                                            300
                        295
Lys Ala Gly Lys Met Tyr Phe Ala Phe Asn Pro Lys Leu Cys Val Ser
                    310
                                        315
Glu Ile Tyr Arg Met Glu Glu Val Thr Gly Thr Lys Gly Arg Gln Ser
                325
                                    330
Lys Gly Asp Ile Asn Thr Arg Asn Asn Gly Glu Arg Ala Ser Cys Glu
                                345
Ser Asp Val Leu His Phe Thr Ser Thr Thr Thr Ser Lys Asn
                            360
                                                365
        355
```

<212> PRT

```
<211> 370
<212> PRT
<213> Homo sapiens
<400> 104
Arg Gly Ser Val Arg Ile Glu Lys Asn Asn Glu Leu Cys Tyr Leu Ala
                                  1.0
Thr Ile Asp Trp Ser Arg Ile Leu Asp Ser Val Glu Asp Asn Tyr Ile
           20
Val Leu Asn Lys Asp Asp Asn Glu Glu Cys Gly Asp Ile Cys Pro Gly
                          40
Thr Ala Lys Gly Lys Thr Asn Cys Pro Ala Thr Val Ile Asn Gly Gln
                       55
                                          60
Phe Val Glu Arg Cys Trp Thr His Ser His Cys Gln Lys Val Cys Pro
                   70
                                      75
Thr Ile Cys Lys Ser His Gly Cys Thr Ala Glu Gly Leu Cys Cys His
              85
                                  90
Ser Glu Cys Leu Gly Asn Cys Ser Gln Pro Asp Asp Pro Thr Lys Cys
                             105
                                                  110
           100
Val Ala Cys Arg Asn Phe Tyr Leu Asp Gly Arg Cys Val Glu Thr Cys
                                             125
       115 120
Pro Pro Pro Tyr Tyr His Phe Gln Asp Trp Arg Cys Val Asn Phe Ser
                      135
                           140
Phe Cys Gln Asp Leu His His Lys Cys Lys Asn Ser Arg Arg Gln Gly
                                     155
                  150
Cys His Gln Tyr Val Ile His Asn Asn Lys Cys Ile Pro Glu Cys Pro
              165 170
                                                      175
Ser Gly Tyr Thr Met Asn Ser Ser Asn Leu Leu Cys Thr Pro Cys Leu
                                                 190
               185
Gly Pro Cys Pro Lys Val Cys His Leu Leu Glu Gly Glu Lys Thr Ile
                         200
                                             205
Asp Ser Val Thr Ser Ala Gln Glu Leu Arg Gly Cys Thr Val Ile Asn
                                          220
                      215
Gly Ser Leu Ile Ile Asn Ile Arg Gly Gly Asn Asn Leu Ala Ala Glu
                   230
                                      235
Leu Glu Ala Asn Leu Gly Leu Ile Glu Glu Ile Ser Gly Tyr Leu Lys
               245
                                  250
Ile Arg Arg Ser Tyr Ala Leu Val Ser Leu Ser Phe Phe Arg Lys Leu
                               265
           260
Arg Leu Ile Arg Gly Glu Thr Leu Glu Ile Gly Asn Tyr Ser Phe Tyr
                           280
                                              285
Ala Leu Asp Asn Gln Asn Leu Arg Gln Leu Trp Asp Trp Ser Lys His
                       295
                                          300
Asn Leu Thr Ile Thr Gln Gly Lys Leu Phe Phe His Tyr Asn Pro Lys
                   310
                                      315
Leu Cys Leu Ser Glu Ile His Lys Met Glu Glu Val Ser Gly Thr Lys
               325
                                  330
Gly Arg Gln Glu Arg Asn Asp Ile Ala Leu Lys Thr Asn Gly Asp Gln
                               345
Ala Ser Cys Glu Asn Glu Leu Leu Lys Phe Ser Tyr Ile Arg Thr Ser
                           360
Phe Asp
    370
<210> 105
<211> 383
```

Ţ

<213> Drosophila melanogaster

```
<400> 105
Arg Gly Gly Val Arg Ile Glu Lys Asn His Lys Leu Cys Tyr Asp Arg
Thr Ile Asp Trp Leu Glu Ile Leu Ala Glu Asn Glu Ser Gln Leu Val
Val Leu Thr Glu Asn Gly Lys Glu Lys Glu Cys Ser Leu Ser Lys Cys
Pro Gly Glu Ile Arg Ile Glu Glu Gly His Asp Asn Thr Ala Ile Glu
                        55
Gly Glu Leu Asn Ala Ser Cys Gln Leu His Asn Asn Arg Arg Leu Cys
                   70
Trp Asn Ser Lys Leu Cys Gln Thr Lys Cys Pro Glu Lys Cys Arg Asn
                                   90
Asn Cys Ile Asp Glu His Thr Cys Cys Ser Gln Asp Cys Leu Gly Gly
                               105
Cys Val Ile Asp Lys Asn Gly Asn Glu Ser Cys Ile Ser Cys Arg Asn
                            120
Val Ser Phe Asn Asn Ile Cys Met Asp Ser Cys Pro Lys Gly Tyr Tyr
                       135
Gln Phe Asp Ser Arg Cys Val Thr Ala Asn Glu Cys Ile Thr Leu Thr
                    150
                                        155
Lys Phe Glu Thr Asn Ser Val Tyr Ser Gly Ile Pro Tyr Asn Gly Gln
               165
                                    170
Cys Ile Thr His Cys Pro Thr Gly Tyr Gln Lys Ser Glu Asn Lys Arg
                                185
                                                    190
           180
Met Cys Glu Pro Cys Pro Gly Gly Lys Cys Asp Lys Glu Cys Ser Ser
                           200
                                                205
Gly Leu Ile Asp Ser Leu Glu Arg Ala Arg Glu Phe His Gly Cys Thr
                                            220
                       215
Ile Ile Thr Gly Thr Glu Pro Leu Thr Ile Ser Ile Lys Arg Glu Ser
                    230
                                        235
Gly Ala His Val Met Asp Glu Leu Lys Tyr Gly Leu Ala Ala Val His
                                   250
               245
Lys Ile Gln Ser Ser Leu Met Val His Leu Thr Tyr Gly Leu Lys Ser
            260
                                265
Leu Lys Phe Phe Gln Ser Leu Thr Glu Ile Ser Gly Asp Pro Pro Met
                                                285
                           280
Asp Ala Asp Lys Tyr Ala Leu Tyr Val Leu Asp Asn Arg Asp Leu Asp
                                            300
                        295
Glu Leu Trp Gly Pro Asn Gln Thr Val Phe Ile Arg Lys Gly Gly Val
                    310
                                        315
Phe Phe His Phe Asn Pro Lys Leu Cys Val Ser Thr Ile Asn Gln Leu
                                    330
Leu Pro Met Leu Ala Ser Lys Pro Lys Phe Phe Glu Lys Ser Asp Glu
                                345
Gly Ala Asp Ser Asn Gly Asn Arg Gly Ser Cys Gly Thr Ala Val Leu
                            360
                                                365
Asn Val Thr Leu Gln Ser Val Gly Ala Asn Ser Ala Ser Leu Asn
    370
```

<210> 106

<211> 381

<212> PRT

<213> Caenorhabditis elegans

```
<400> 106
Asn Gly Gly Val Arg Ile Ile Asp Asn Arg Lys Leu Cys Tyr Thr Lys
                                  10
Thr Ile Asp Trp Lys His Leu Ile Thr Ser Ser Ile Asn Asp Val Val
Val Asp Asn Ala Ala Glu Tyr Ala Val Thr Glu Thr Gly Leu Met Cys
Pro Arg Gly Ala Cys Glu Glu Asp Lys Gly Glu Ser Lys Cys His Tyr
Leu Glu Glu Lys Asn Gln Glu Gln Gly Val Glu Arg Val Gln Ser Cys
                   70
Trp Ser Asn Thr Thr Cys Gln Lys Ser Cys Ala Tyr Asp Arg Leu Leu
               85
                                   90
Pro Thr Lys Glu Ile Gly Pro Gly Cys Asp Ala Asn Gly Asp Arg Cys
                               105
His Asp Gln Cys Val Gly Gly Cys Glu Arg Val Asn Asp Ala Thr Ala
                           120
Cys His Ala Cys Lys Asn Val Tyr His Lys Gly Lys Cys Ile Glu Lys
                       135
                                           140
Cys Asp Ala His Leu Tyr Leu Leu Leu Gln Arg Arg Cys Val Thr Arg
                                       155
                   150
Glu Gln Cys Leu Gln Leu Asn Pro Val Leu Ser Asn Lys Thr Val Pro
               165
                                   170
Ile Lys Ala Thr Ala Gly Leu Cys Ser Asp Lys Cys Pro Asp Gly Tyr
                              185
                                                   190
          180
Gln Ile Asn Pro Asp Asp His Arg Glu Cys Arg Lys Cys Val Gly Lys
                                               205
                           200
Cys Glu Ile Val Cys Glu Ile Asn His Val Ile Asp Thr Phe Pro Lys
                                           220
                       215
Ala Gln Ala Ile Arg Leu Cys Asn Ile Ile Asp Gly Asn Leu Thr Ile
                                       235
                   230
Glu Ile Arg Gly Lys Gln Asp Ser Gly Met Ala Ser Glu Leu Lys Asp
                                   250
               245
Ile Phe Ala Asn Ile His Thr Ile Thr Gly Tyr Leu Leu Val Arg Gln
                               265
           260
Ser Ser Pro Phe Ile Ser Leu Asn Met Phe Arg Asn Leu Arg Arg Ile
                                               285
                           280
Glu Ala Lys Ser Leu Phe Arg Asn Leu Tyr Ala Ile Thr Val Phe Glu
                      295
                                           300
Asn Pro Asn Leu Lys Lys Leu Phe Asp Ser Thr Thr Asp Leu Thr Leu
                   310
                                       315
Asp Arg Gly Thr Val Ser Ile Ala Asn Asn Lys Met Leu Cys Phe Lys
               325
                                  330
Tyr Ile Lys Gln Leu Met Ser Lys Leu Asn Ile Pro Leu Asp Pro Ile
                               345
Asp Gln Ser Glu Gly Thr Asn Gly Glu Lys Ala Ile Cys Glu Asp Met
                           360
Ala Ile Asn Val Ser Ile Thr Ala Val Asn Ala Asp Ser
                        375
```

<211> 370

<212> PRT

<213> Homo sapiens

<400> 107

Ala Leu Pro Val Ala Val Leu Leu Ile Val Gly Gly Leu Val Ile Met

```
10
Leu Tyr Val Phe His Arg Lys Arg Asn Asn Ser Arg Leu Gly Asn Gly
Val Leu Tyr Ala Ser Val Asn Pro Glu Tyr Phe Ser Ala Ala Asp Val
                           40
Tyr Val Pro Asp Glu Trp Glu Val Ala Arg Glu Lys Ile Thr Met Ser
                       55
Arg Glu Leu Gly Gln Gly Ser Phe Gly Met Val Tyr Glu Gly Val Ala
                   70
Lys Gly Val Val Lys Asp Glu Pro Glu Thr Arg Val Ala Ile Lys Thr
                                  90
Val Asn Glu Ala Ala Ser Met Arg Glu Arg Ile Glu Phe Leu Asn Glu
                              105
Ala Ser Val Met Lys Glu Phe Asn Cys His His Val Val Arg Leu Leu
                           120
       115
Gly Val Val Ser Gln Gly Gln Pro Thr Leu Val Ile Met Glu Leu Met
                       135
Thr Arg Gly Asp Leu Lys Ser Tyr Leu Arg Ser Leu Arg Pro Glu Met
                  150
                                       155
Glu Asn Asn Pro Val Leu Ala Pro Pro Ser Leu Ser Lys Met Ile Gln
               165
                                   170
                                                       175
Met Ala Gly Glu Ile Ala Asp Gly Met Ala Tyr Leu Asn Ala Asn Lys
                               185
                                                  190
           180
Phe Val His Arg Asp Leu Ala Ala Arg Asn Cys Met Val Ala Glu Asp
       195
                          200
                                              205
Phe Thr Val Lys Ile Gly Asp Phe Gly Met Thr Arg Asp Ile Tyr Glu
                       215
                                           220
Thr Asp Tyr Tyr Arg Lys Gly Gly Lys Gly Leu Leu Pro Val Arg Trp
                                       235
                   230
Met Ser Pro Glu Ser Leu Lys Asp Gly Val Phe Thr Thr Tyr Ser Asp
                                  250
               245
Val Trp Ser Phe Gly Val Val Leu Trp Glu Ile Ala Thr Leu Ala Glu
                              265
Gln Pro Tyr Gln Gly Leu Ser Asn Glu Gln Val Leu Arg Phe Val Met
                           280
Glu Gly Gly Leu Leu Asp Lys Pro Asp Asn Cys Pro Asp Met Leu Phe
                       295
                                           300
Glu Leu Met Arg Met Cys Trp Gln Tyr Asn Pro Lys Met Arg Pro Ser
                   310
                                      315
Phe Leu Glu Ile Ile Ser Ser Ile Lys Glu Met Glu Pro Gly Phe
                                   330
Arg Glu Val Ser Phe Tyr Tyr Ser Glu Glu Asn Lys Leu Pro Glu Pro
                               345
Glu Glu Leu Asp Leu Glu Pro Glu Asn Met Glu Ser Val Pro Leu Asp
                           360
Pro Ser
    370
<210> 108
<211> 374
<212> PRT
<213> Homo sapiens
<400> 108
Ile Gly Pro Leu Ile Phe Val Phe Leu Phe Ser Val Val Ile Gly Ser
                                   10
```

Ile Tyr Leu Phe Leu Arg Lys Arg Gln Pro Asp Gly Pro Leu Gly Pro

```
20
                                25
Leu Tyr Ala Ser Ser Asn Pro Glu Tyr Leu Ser Ala Ser Asp Val Phe
Pro Cys Ser Val Tyr Val Pro Asp Glu Trp Glu Val Ser Arg Glu Lys
                        55
Ile Thr Leu Leu Arg Glu Leu Gly Gln Gly Ser Phe Gly Met Val Tyr
Glu Gly Asn Ala Arg Asp Ile Ile Lys Gly Glu Ala Glu Thr Arg Val
                                   90
Ala Val Lys Thr Val Asn Glu Ser Ala Ser Leu Arg Glu Arg Ile Glu
                                105
Phe Leu Asn Glu Ala Ser Val Met Lys Gly Phe Thr Cys His His Val
                            120
Val Arg Leu Leu Gly Val Val Ser Lys Gly Gln Pro Thr Leu Val Val
                                            140
                        135
Met Glu Leu Met Ala His Gly Asp Leu Lys Ser Tyr Leu Arg Ser Leu
                   150
                                        155
Arg Pro Glu Ala Glu Asn Asn Pro Gly Arg Pro Pro Pro Thr Leu Gln
                                    170
               165
Glu Met Ile Gln Met Ala Ala Glu Ile Ala Asp Gly Met Ala Tyr Leu
                                185
           180
Asn Ala Lys Lys Phe Val His Arg Asp Leu Ala Ala Arg Asn Cys Met
                            200
Val Ala His Asp Phe Thr Val Lys Ile Gly Asp Phe Gly Met Thr Arg
                        215
                                            220
Asp Ile Tyr Glu Thr Asp Tyr Tyr Arg Lys Gly Gly Lys Gly Leu Leu
                    230
                                        235
Pro Val Arg Trp Met Ala Pro Glu Ser Leu Lys Asp Gly Val Phe Thr
                                    250
               245
Thr Ser Ser Asp Met Trp Ser Phe Gly Val Val Leu Trp Glu Ile Thr
                                265
                                                   270
Ser Leu Ala Glu Gln Pro Tyr Gln Gly Leu Ser Asn Glu Gln Val Leu
                                                285
                            280
Lys Phe Val Met Asp Gly Gly Tyr Leu Asp Gln Pro Asp Asn Cys Pro
                                            300
                        295
Glu Arg Val Thr Asp Leu Met Arg Met Cys Trp Gln Phe Asn Pro Lys
                                        315
                    310
Met Arg Pro Thr Phe Leu Glu Ile Val Asn Leu Leu Lys Asp Asp Leu
                                    330
His Pro Ser Phe Pro Glu Val Ser Phe Phe His Ser Glu Glu Asn Lys
                                345
Ala Pro Glu Ser Glu Glu Leu Glu Met Glu Phe Glu Asp Met Glu Asn
                            360
Val Pro Leu Asp Arg Ser
    370
```

<211> 384

<212> PRT

<213> Drosophila melanogaster

<400> 109

Gly Ile Gly Leu Ala Phe Leu Ile Val Ser Leu Phe Gly Tyr Val Cys 1 5 10 15

Tyr Leu His Lys Arg Lys Val Pro Ser Asn Asp Leu His Met Asn Thr 20 25 30

Glu Val Asn Pro Phe Tyr Ala Ser Met Gln Tyr Ile Pro Asp Asp Trp

```
Glu Val Leu Arg Glu Asn Ile Ile Gln Leu Ala Pro Leu Gly Gln Gly
Ser Phe Gly Met Val Tyr Glu Gly Ile Leu Lys Ser Phe Pro Pro Asn
                    70
Gly Val Asp Arg Glu Cys Ala Ile Lys Thr Val Asn Glu Asn Ala Thr
                                    90
Asp Arg Glu Arg Thr Asn Phe Leu Ser Glu Ala Ser Val Met Lys Glu
                                105
Phe Asp Thr Tyr His Val Val Arg Leu Leu Gly Val Cys Ser Arg Gly
                            120
Gln Pro Ala Leu Val Val Met Glu Leu Met Lys Lys Gly Asp Leu Lys
                        135
Ser Tyr Leu Arg Ala His Arg Pro Glu Glu Arg Asp Glu Ala Met Met
                    150
                                        155
Thr Tyr Leu Asn Arg Ile Gly Val Thr Gly Asn Val Gln Pro Pro Thr
                                    170
                165
Tyr Gly Arg Ile Tyr Gln Met Ala Ile Glu Ile Ala Asp Gly Met Ala
           180
                                185
                                                    190
Tyr Leu Ala Ala Lys Lys Phe Val His Arg Asp Leu Ala Ala Arg Asn
       195
                            200
                                                205
Cys Met Val Ala Asp Asp Leu Thr Val Lys Ile Gly Asp Phe Gly Met
                        215
                                            220
Thr Arg Asp Ile Tyr Glu Thr Asp Tyr Tyr Arg Lys Gly Thr Lys Gly
                    230
                                        235
Leu Leu Pro Val Arg Trp Met Pro Pro Glu Ser Leu Arg Asp Gly Val
                245
                                    250
Tyr Ser Ser Ala Ser Asp Val Phe Ser Phe Gly Val Val Leu Trp Glu
                                265
Met Ala Thr Leu Ala Ala Gln Pro Tyr Gln Gly Leu Ser Asn Glu Gln
                            280
                                                285
Val Leu Arg Tyr Val Ile Asp Gly Gly Val Met Glu Arg Pro Glu Asn
                        295
                                            300
Cys Pro Asp Phe Leu His Lys Leu Met Gln Arg Cys Trp His His Arg
                                        315
                    310
Ser Ser Ala Arg Pro Ser Phe Leu Asp Ile Ile Ala Tyr Leu Glu Pro
                325
                                    330
Gln Cys Pro Asn Ser Gln Phe Lys Glu Val Ser Phe Tyr His Ser Glu
                                345
            340
Ala Gly Leu Gln His Arg Glu Lys Glu Arg Lys Glu Arg Asn Gln Leu
                            360
                                                365
Asp Ala Phe Ala Ala Val Pro Leu Asp Gln Asp Leu Gln Asp Arg Glu
    370
                        375
```

<211> 380

<212> PRT

<213> Caenorhabditis elegans

<400> 110

```
55
                                            60
Gln Cys Gly Glu Gly Ser Phe Gly Lys Val Tyr Leu Gly Thr Gly Asn
                    70
                                       75
Asn Val Val Ser Leu Met Gly Asp Arg Phe Gly Pro Cys Ala Ile Lys
Ile Asn Val Asp Asp Pro Ala Ser Thr Glu Asn Leu Asn Tyr Leu Met
                               105
                                                   110
Glu Ala Asn Ile Met Lys Asn Phe Lys Thr Asn Phe Ile Val Gln Leu
                            120
Tyr Gly Val Ile Ser Thr Val Gln Pro Ala Met Val Val Met Glu Met
                                            140
                        135
Met Asp Leu Gly Asn Leu Arg Asp Tyr Leu Arg Ser Lys Arg Glu Asp
                    150
                                        155
Glu Val Phe Asn Glu Thr Asp Cys Asn Phe Phe Asp Ile Ile Pro Arg
                                    170
                165
Asp Lys Phe His Glu Trp Ala Ala Gln Ile Cys Asp Gly Met Ala Tyr
           180
                                185
Leu Glu Ser Leu Lys Phe Cys His Arg Asp Leu Ala Ala Arg Asn Cys
                            200
                                                205
       195
Met Ile Asn Arg Asp Glu Thr Val Lys Ile Gly Asp Phe Gly Met Ala
                        215
                                            220
Arg Asp Leu Phe Tyr His Asp Tyr Tyr Lys Pro Ser Gly Lys Arg Met
                   230
                                        235
Met Pro Val Arg Trp Met Ser Pro Glu Ser Leu Lys Asp Gly Lys Phe
                245
                                    250
Asp Ser Lys Ser Asp Val Trp Ser Phe Gly Val Val Leu Tyr Glu Met
                                                    270
                                265
            260
Val Thr Leu Gly Ala Gln Pro Tyr Ile Gly Leu Ser Asn Asp Glu Val
                            280
                                                285
Leu Asn Tyr Ile Gly Met Ala Arg Lys Val Ile Lys Lys Pro Glu Cys
                        295
                                            300
Cys Glu Asn Tyr Trp Tyr Lys Val Met Lys Met Cys Trp Arg Tyr Ser
                                        315
                    310
Pro Arg Asp Arg Pro Thr Phe Leu Gln Leu Val His Leu Leu Ala Ala
                                   330
                325
Glu Ala Ser Pro Glu Phe Arg Asp Leu Ser Phe Val Leu Thr Asp Asn
            340
                                345
Gln Met Ile Leu Asp Asp Ser Glu Ala Leu Asp Leu Asp Asp Ile Asp
                            360
Asp Thr Asp Met Asn Asp Gln Val Val Glu Val Ala
                        375
```

<210> 111 <211> 103

<212> PRT

<213> Caenorhabditis elegans

<400> 111

```
70
                                        75
65
Arg His Val Asp His Cys Lys His Ala Phe Glu Met Lys Ser Asp Met
                                    90
               85
Val Cys Val Asn Pro Tyr His
            100
<210> 112
<211> 104
<212> PRT
<213> Homo sapiens
<400> 112
Gly Gly Glu Ser Glu Thr Phe Ala Lys Arg Ala Ile Glu Ser Leu Val
Lys Lys Leu Lys Glu Lys Lys Asp Glu Leu Asp Ser Leu Ile Thr Ala
Ile Thr Thr Asn Gly Ala His Pro Ser Lys Cys Val Thr Ile Gln Arg
Thr Leu Asp Gly Arg Leu Gln Val Ala Gly Arg Lys Gly Phe Pro His
                        55
Val Ile Tyr Ala Arg Leu Trp Arg Trp Pro Asp Leu His Lys Asn Glu
                                         75
                    70
Leu Lys His Val Lys Tyr Cys Gln Tyr Ala Phe Asp Leu Lys Cys Asp
                85
Ser Val Cys Val Asn Pro Tyr His
            100
<210> 113
<211> 205
<212> PRT
<213> Caenorhabditis elegans
<400> 113
Ile Val Tyr Tyr Glu Lys Asn Leu Gln Ile Gly Glu Lys Lys Cys Ser
                                     10
Arg Gly Asn Phe His Val Asp Gly Gly Phe Ile Cys Ser Glu Asn Arg
                                 25
 Tyr Ser Leu Gly Leu Glu Pro Asn Pro Ile Arg Glu Pro Val Ala Phe
                             40
 Lys Val Arg Lys Ala Ile Val Asp Gly Ile Arg Phe Ser Tyr Lys Lys
                         55
                                             60
 Asp Gly Ser Val Trp Leu Gln Asn Arg Met Lys Tyr Pro Val Phe Val
                                         75
                     70
 Thr Ser Gly Tyr Leu Asp Glu Gln Ser Gly Gly Leu Lys Lys Asp Lys
                                                          95
                                     90
 Val His Lys Val Tyr Gly Cys Ala Ser Ile Lys Thr Phe Gly Phe Asn
                                                      110
                                 105
             100
 Val Ser Lys Gln Ile Ile Arg Asp Ala Leu Leu Ser Lys Gln Met Ala
                                                 125
                             120
 Thr Met Tyr Leu Gln Gly Lys Leu Thr Pro Met Asn Tyr Ile Tyr Glu
                         135
 Lys Lys Thr Gln Glu Glu Leu Arg Arg Glu Ala Thr Arg Thr Thr Asp
                                         155
                     150
 Ser Leu Ala Lys Tyr Cys Cys Val Arg Val Ser Phe Cys Lys Gly Phe
                                      170
```

Gly Glu Ala Tyr Pro Glu Arg Pro Ser Ile His Asp Cys Pro Val Trp

```
180
                            185
Ile Glu Leu Lys Ile Asn Ile Ala Tyr Asp Phe Met Asp
            200
<210> 114
<211> 212
<212> PRT
<213> Homo sapiens
<400> 114
Ile Ala Tyr Phe Glu Met Asp Val Gln Val Gly Glu Thr Phe Lys Val
              5
                                10
       20
                             25
                        40
                     55
```

Pro Ser Ser Cys Pro Ile Val Thr Val Asp Gly Tyr Val Asp Pro Ser Gly Gly Asp Arg Phe Cys Leu Gly Gln Leu Ser Asn Val His Arg Thr Glu Ala Ile Glu Arg Ala Arg Leu His Ile Gly Lys Gly Val Gln Leu Glu Cys Lys Gly Glu Gly Asp Val Trp Val Arg Cys Leu Ser Asp His 70 75 Ala Val Phe Val Gln Ser Tyr Tyr Leu Asp Arg Glu Ala Gly Arg Ala 85 90 Pro Gly Asp Ala Val His Lys Ile Tyr Pro Ser Ala Tyr Ile Lys Val 100 105 110 Phe Asp Leu Arg Gln Cys His Arg Gln Met Gln Gln Gln Ala Ala Thr 115 120 125 Ala Gln Ala Ala Ala Ala Gln Ala Ala Ala Val Ala Gly Asn Ile 135 140 Pro Gly Pro Gly Ser Val Gly Gly Ile Ala Pro Ala Ile Ser Leu Ser 150 155 Ala Ala Ala Gly Ile Gly Val Asp Asp Leu Arg Arg Leu Cys Ile Leu 170 175 165 Arg Met Ser Phe Val Lys Gly Trp Gly Pro Asp Tyr Pro Arg Gln Ser 180 185 190 Ile Lys Glu Thr Pro Cys Trp Ile Glu Ile His Leu His Arg Ala Leu 195 200 Gln Leu Leu Asp